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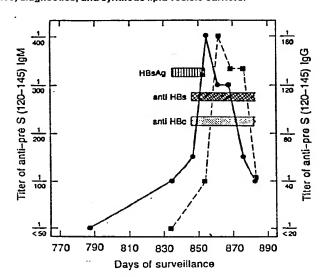
#### **EUROPEAN PATENT APPLICATION**

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- Pre-S gene coded peptide hepatitis B immunogens, vaccines, diagnostics, and synthetic lipid vesicle carriers.
- A hepatitis B vaccine containing a peptide with an amino acid chain of at least six consecutive amino acids within the pre-S gene coded region of the envelope of hepatitis B virus. The vaccine being free of an amino acid sequence corresponding to the naturally occuring evelope proteins of hepatitis B virus and a physiologically acceptable diluent. The peptide being free or linked to a carrier. The carrier being a conventional carrier or a novel carrier including a lipid vesicle stabilized by cross-linking and having covalently bonded active sites on the outer surface thereon. Such novel carrier being useful not only to link the novel peptide containing an amino acid chain with amino acids within the pre-S gene coded region of the surface antigen of hepatitis B virus, but can also be used to bind synthetic peptide analogues of other viral proteins, as well as bacterial, allergen and parasitic proteins of man and animals. The peptides of the invention can be utilized in diagnostics for the detection of antigens and antibodies.



#### BACKGROUND OF THE INVENTION

The present invention concerns pre-S gene coded hepatitis B immunogens, vaccines and diagnostics. More especially, this invention concerns novel pre-S gene coded peptides and novel carriers, particularly carriers for pre-S gene coded peptides. Even more especially, the present invention relates to synthetic pre-S gene coded peptides covalently linked to lipid vesicle carriers.

There are approximately 600,000 persistent carriers of hepatitis B virus (HBV) in the United States; the estimated total number of carriers in the world is 200 million. A considerable portion of HBV carriers have chronic liver disease. The involvement of HBV in liver cancer has been demonstrated (W. Szmuness, Prog. Med. Virol. 24, 40 (1978) and R.P. Beasley, L.-Y. Hwang, C.-C. Ling, C.-S. Chien, Lancet Nov., 21, 1129 (1981)).

HBV infections thus represent a major public health problem worldwide. Already available vaccines (S. Krugman, in <u>Viral Hepatitis: Laboratory and Clinical Science</u>, F. Deinhardt, J. Deinhardt, Eds., Marcel Dekker, Inc., New York-Basel, 1983, pp. 257-263) produced from the serum of HBV carriers, because of limited resources and production costs involved, do not provide the appropriate

1 means to control and eradicate the disease worldwide. 2 is hope, however, that this may be accomplished by vacci. 3 based on recombinant DNA technology and/or synthetic 4 peptides. 5 The biology, structure and immunochemistry of HB. 6 and the genetic organization of its DNA genome have been 7 reviewed (B.S. Blumberg, Science, 197 17, (1977)). 8 cloning and sequencing of the genome of several hepatitis 9 virus (HBV) isolates led to the elucidation of the genetic 10 structure of the viral DNA (P. Tiollais, P. Charnay, G.N. 11 Vyas, Science, 213, 406, (1981)). 12 The immunologic markers of HBV infection include 13 the surface antigen (HBsAg), the core antigen (HBcAg), the 14 "e" antigen (HBeAg) and their respective antibodies. 15 Antibodies against HBsAg are protective against HBV 16 infection. 17 Several antigenic subtypes of HBV and of subvira 18 approximately 22 nm diameter particles (hepatitis B surfac 19 antigen; HBsAg) have been recognized (G. Le Bouvier, A. 20 Williams, Am. J. Med. Sci., 270, 165 (1975)). All of thes 21 subtypes (for example, ayw, adyw, adw2, adw and adr) share 22 common (group-specific) envelope epitopes, the immune 23 response against which appears sufficient for protection 24 against infection by any of the virus subtypes (W. Szmunes 25 C.E. Stevens, E.J. Harley, E.A. Zang, H.J. Alter, P.E. 26 Taylor, A. DeVera, G.T.S. Chen, A. Kellner, et al., N. Eng 27 J. Med., 307, 1481, (1982)). 28

The physical structure and proposed genetic

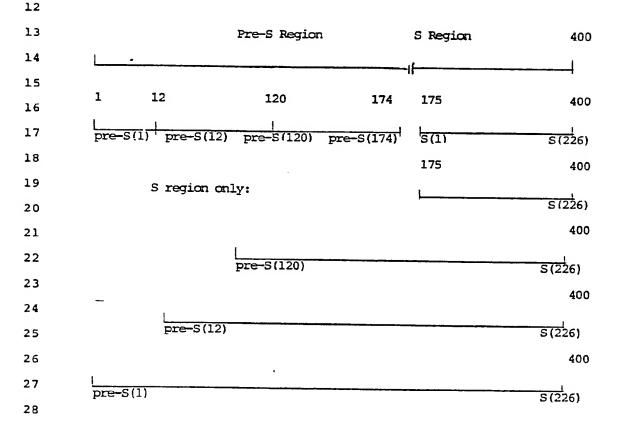
organization of the HBV genome are described by Tiollais  $\epsilon$ 

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al, 1981, <u>supra</u> at p<sup>-</sup> 408-409. There are two DNA strands, namely the long (L) strand and the short (S) strand. The L strand transcript has four open reading frame regions which are termed (S + pre-S), C, P and X.

The open reading frame region (S + pre-S) corresponds to the envelope (env) gene of HBV DNA and codes for a family of proteins found in the HBV envelope and in virus related particles.

A schematic representation of the potential translation products of the env gene(s) of HBV DNA is as follows:



The numbers in the above schematic refers to amino acids (AA). A translation initiation site at Met 1 exists

for the adw<sub>2</sub> and adr substypes onl,. The first amino acid for the other subtypes correspond to position pre-S 12.

Hereinafter, amino acid sequences corresponding to the pre-S region (env 1 to 174) are designated with the prefix "pre-S" and amino acid sequences corresponding to the S region (env 175 to 400) are designated by the prefix "S". In the env gene product representation, the S region spans amino acids 175 to 400 as compared to amino acids 1 to 226 in the "S region only" representation.

In the above schematic, the pre-S region is defined by amino acid sequence positions pre-S 1 to amino acid sequence position pre-S 174. The S region is defined by sequence positions S 1 (amino acid 175 of the open reading frame and adjacent to pre-S 174) to sequence position S 266 (amino acid 400 of the open reading frame). The s-gene product (S-protein) consists of this 226 amino acid sequence.

The epitope(s) essential for eliciting virus-neutralizing antibodies have not yet been unambiguously defined. It has been reported that the group-specificity is represented by a complex of determinants located on each of the two major approximately 22 and approximately 26 kilodalton constituent proteins (P22 and P26) of the virus envelope and of the hepatitis B surface antigen (HBsAg). See J.W.-K. Shih, J.L. Gerin, J. Immunol., 115, 634, (1975); J.W.-K. Shih, P.L. Tan, J.L. Gerin, J. Immunol., 120, 520, (1978); S. Mishiro, M. Imai, K. Takahashi, A. Machida, T. Gotanda, Y. Miyakawa, M.

Mayumi, J. Immunol., 124, 1589, (1980); and G.R. Dreesman, 1 R. Chairez, M. Suarez., F.B. Hollinger, R.J. Courtney, J.L. Melnick, J. Virol., 16, 508, (1975). 3 These proteins have identical amino acid sequences coded for by the S-gene of HBV DNA (Tiollais et al, supra), but the larger protein also carries carbohydrate chains. Peptides corresponding to selected segments of the S-gene product were synthesized and shown to elicit antibodies to HBsAg (anti-HBs). However, immunization of chimpanzees with these peptides resulted in only partial protection against 10 11 HBV infection (N. Williams, Nature, 306, 427, (1983)). 12 It has been reported recently that the minor 13 glycoprotein components of HBsAg with  $M_r$  of approximately 33 14 and approximately 36 kilodaltons (P33, P36) are coded for 15 HBV DNA and contain the sequence of P22 (226 amino acids 16 corresponding to the S region) and have 55 additional amino 17 acids at the amino-terminal part which are coded by the 18 pre-S region of the env gene(s) of HBV DNA. See W. Stibbe, 19 W.H. Gerlich, <u>Virology</u>, <u>123</u>, 436, (1982); M.A. Feitelson, 20 P.L. Marion, W.S. Robinson, <u>Virology</u>, <u>130</u>, 76, (1983); W. 21 Stibbe, W.H. Gerlich, J. Virol., 46, 626, (1983); and A. 22 Machida, S. Kishimoto, H. Ohnuma, H. Miyamoto, K. Baba, K. 23 Oda, T. Nakamura, Y. Miyakawa, M. Mayumi, Gastroenterology, 24 85, 268, (1983). Machida et al describe an amino acid 25 sequence composition as a receptor for polymerized serum 26 albumin. 27 Heretofore, amino acid sequences coded for by the 28 pre-S region of the hepatitis B virus DNA were virtually 29

completely ignored for purposes of producing synthetic vaccines. The hepatitis B vaccine currently in use in the United States lacks the pre-S gene coded sequences (and therefore does not elicit antibodies to such sequences) and thus elicits an immune response to the HBV envelope which is incomplete as compared with that occurring during recovery from natural infection.

The generation of antibodies to proteins by immunization with short peptides having the amino acid sequence corresponding to the sequence of preselected protein fragments appears to be a frequent event (Nima, H.L., Houghten, R.A., Walker, L.E., Reisfeld, R.A., Wilson, I.A., Hogle, J.M. and Lerner, R.A., "Generation Of Protein-Reactive Antibodies By Short Peptides Is An Event Of High Frequency: Implications For The Structural Basis Of Immune Recognition", Proceedings of the National Academy of Sciences USA, 80, 4949-4953, (1983)). Nevertheless, the generation of antibodies which recognize the native protein may depend on the appropriate conformation of the synthetic peptide immunogen and on other factors not yet understood. See Pfaff, E., Mussgay, M., Böhm, H.O., Schulz, G.E. and Schaller, H., "Antibodies Against A Preselected Peptide Recognize And Neutralize Foot And Mouth Disease Virus", The EMBO Journal, 7, 869-874, (1982); Neurath, A.R., Kent, S.B.H. and Strick, N., "Specificity Of Antibodies Elicited By A Synthetic Peptide Having A Sequence In Common With A

Fragment Of A Virus Protein, The Hepatitis B Surface 1 Antigen," Proceedings Of The National Academy Of Sciences USA, 79, 7871-7875, (1982); Ionescu-Matiu, J., Kennedy, R.C., Sparrow, J.T., Culwell, A.R., Sanchez, Y., Melnick, J.L. and Dreesman, G.R., "Epitopes Associated With A 6 Synthetic Hepatitis B Surface Antigen Peptide", The Journal Of Immunology, 130, 1947-1952, (1983); and Kennedy, R.C., Dreesman, G.R., Sparrow, J.T., Culwell, A.R., Sanchez, Y., 9 Ionescu-Matiu, I., Hollinger, F.B. and Melnick, J.L. (1983); 10 "Inhibition Of A Common Human Anti-Hepatitis B Surface 11 Antigen Idiotype By A Cyclic Synthetic Peptide," Journal of 12 Virology, 46, 653-655, (1983). For this reason, immunization 13 with synthetic peptide analogues of various virus proteins 14 has only rarely resulted in production of virus-neutralizing 15 antisera comparable to those elicited by the viruses (virus 16 proteins) themselves (Pfaff et al., 1982, supra). Thus, the 17 preparation of synthetic immunogens optimally mimicking 18 antigenic determinants on intact viruses remains a 19 challenge. 20 Replacement of commonly used protein carriers, 21 namely keyhole limpet hemocyanin (KLH), albumin, etc., by 22 synthetic carriers, represents part of such challenge. 23 Although recent reports indicate that free synthetic 24 peptides can be immunogenic, (Dreesman, G.R., Sanchez, Y., 25 Ionescu-Matiu, I., Sparrow, J.T., Six, H.R., Peterson, D.L., 26 Hollinger, F.B. and Melnick, J.L., "Antibody To Hepatitis B 27 28 29

1 pep'ides can be immunogenic, (Dreesmar, G.R., Sanchez, Y., 2 Ionesc -- Matiu, I., Spa\_row, J.T., Six, H.R., Peterson, D.L., 3 Hollinger, F.B. and Melnick, J.L., "Antibody To Hepatitis B 4 Surface Antigen After A Single Inoculation Of Uncoupled 5 Synthetic HBsAg Peptides" Nature, 295, 158-160, (1982), and 6 Schmitz, H.E., Atassi, H., and Atassi, M.Z., "Production Of 7 Monoclonal Antibodies To Surface Regions That Are Non-Immunogenic In A Protein Using Free Synthetic Peptide As Immunogens: Demonstration With Sperm-whale Myoglobin", 10 Immunological Communications, 12, 161-175, (1983)), even in 11 these cases the antibody response was enhanced by linking of 12 the peptides to a protein carrier (Sanchez, Y., 13 Ionescu-Matiu, I., Sparrow, J.T., Melnick, J.L., Dreesman, 14 G.R., "Immunogenicity Of Conjugates And Micelles Of 15 Synthetic Hepatitis B Surface Antigen Peptides", 16 Intervirology, 18, 209-213, (1982)). 17 For commonly used protein carriers there is a 18 strong immune response to the carrier, as well as the 19 synthetic peptide. Thus, it would be advantageous to evoke 20 an anti-HBs response with peptides by use of non-protein 21 carriers, which themselves do not evoke an antibody 22 response. 23 The possible use of several distinct vaccines in 24

prophylaxis would be facilitated by the availability of fully synthetic immunogens.

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                                     DEFINITIONS
 2
         Amino Acid Code Words (as appearing in Fig. 2)
 3
         D
               Asp
                          aspartic acid
 4
         N
               Asn
                          asparagine
 5
         T
               Thr
                          threonine
 6
         S ,
              Ser
                          serine
 7
         E
              Glu
                          glutamic acid
 8
         Q
              Gln
                          glutamine
 9
         P
              Pro
                         proline
10
         G
              Gly
                         glycine
11
         Α
              Ala
                         alanine
12
         С
              Cys
                         cysteine
13
         v
              Val
                         valine
14
         М
              Met
                         methionine
15
         I
              Ile
                         isoleucine
16
         L
              Leu
                         leucine
17
         Y
              Tyr
                         tyrosine
18
         F
              Phe
                         phenylalanine
19
        W
              Trp
                         tryptophane
20
        K
              Lys
                         lysine
21
        Н
              His
                         histidine
22
        R
              Arg
                         arginine
23
        <u>HBV</u>
                         hepatitis B virus
24
        HBsAg
                         hepatitis B surface antigen.
25
        DNA
                         deoxyribonucleic acid
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#### SUMMARY OF THE INVENTION

The applicants have found that antibodies to the pre-S protein appear early in the course of hepatitis B infection and probably play the role of antibodies eliminating HBV from the circulation and thus interrupting further spread of the infection. Antibodies to the pre-S protein are likely to represent virus-neutralizing antibodies. The failure of some hepatitis B vaccines to elicit such antibodies may be of considerable biological significance, as indicated by poor immunoprophylactic effects elicited by such vaccines in some populations, despite a detectable immune response to the S-protein.

Applicants have discovered that amino acid sequences coded for by the pre-S region of the env gene of hepatitis B virus (HBV) DNA carry dominant antigenic determinants common to intact and denatured HBsAg.

Applicants have found that immuno-dominant disulfide bond-independent epitopes recognized by human antibodies to hepatitis B virus (HBV) exist within proteins containing amino acid sequences coded by the pre-S region of HBV DNA, and more particularly within proteins containing an N-terminal portion (coded for the pre-S region of HBV DNA) having an N-terminal methionine at amino acid sequence position pre-S 120. Applicants further discovered that peptides corresponding to amino acid sequences in the pre-S region, and more particularly in the aforementioned region

starting at amino acid 120 of the env gene eper reading.

frame, inhibit the reaction between human anti-HBs and P33 (P36), are highly immunogenic, and elicit high levels of group-specific antibodies against HBsAg and HBV. The immunogenicity of such peptides is enhanced by covalent linking to novel lipid vesicle (liposome) carriers also discovered by applicants.

Glutaraldehyde-fixed liposomes were found by applicants to be preferred carriers for the peptides of the invention for inducing anti-HBs.

The present invention thus concerns a hepatitis B peptide immunogen including a peptide containing an amino acid chain corresponding to at least six consecutive amino acids within the pre-S gene coded region of the envelope of HBV. The hepatitis B peptide immunogen being free of an amino acid chain corresponding to the naturally occurring envelope proteins of hepatitis B virus.

The naturally occurring envelope proteins of hepatitis B virus include the following:

- (1) a full length translational product of the env gene of HBV, i.e., for  $adw_2$  and adr pre-S(1-174) + S(175-400)=400 amino acids, for ayw, adyw and adw pre-S(12-174) + S(1-226) = 389 amino acids (env 12-400);
- (2) pre-S(120-174) + S(175-400) = 281 amino acids (env 120-400) = terminal 55 amino acids in the pre-S region

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+ 226 amino acids comprising the entire S region (the corresponding proteins approximately 33 and 36 kD in size (P33 and P36), and differing from each other in the extent of glycosylation); and

(3) S(1-226) = 226 amino acids, i.e., the entire S region (env 175-400); representing the approximately 22 and 26 kD major constituents of the HBV envelope (P22 and P26) in their non-glycosylated and glycosylated forms (the "S-protein").

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In an embodiment of the hepatitis B peptide immunogen of the present invention, the corresponding chain of amino acids lies between the sequence positions pre-S 120 and pre-S 174. In another embodiment of the invention, the chain of amino acids is between sequence positions pre-S 1 and pre-S 120. In a further embodiment of the invention, the corresponding chain of amino acids includes the methionine amino acid at sequence position pre-S 120. In still another embodiment, the chain of amino acids is an amino acid chain containing at least 26 amino acids in the pre-S region. Still further, the chain of amino acids containing at least 26 amino acids can correspond to a chain of at least 26 consecutive amino acids disposed between sequence position pre-S 120 and sequence position pre-S 174. Generally the peptide has no more than 280 amino acids, preferably no more than 225 amino acids, more preferably no

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more than 174 amino acia, even more preferably no more than 100 amino acids, and still more preferably, no more than 50 amino acids. The vaccine of the present invention can be composed solely of a peptide, or preferably of a peptide joined to a carrier. Such carrier can be a conventional carrier, or a novel carrier according to the present invention as described hereinbelow.

The hepatitis B peptide immunogen of the present invention is free of any serum proteins, e.g., blood serum proteins.

The present invention also concerns a hepatitis B vaccine including a peptide containing an amino acid chain corresponding to at least six consecutive amino acids within the pre-S gene coded region of the envelope of HBV, and a physiologically acceptable diluent, e.g., phosphate buffered saline. The hepatitis B peptide vaccine being free of an amino acid chain corresponding to the naturally occurring envelope proteins of hepatitis B virus.

The present invention is also directed to a novel carrier for peptides. In a particularly preferred embodiment of the present invention, the hepatitis B vaccine containing an amino acid chain corresponding to a chain of amino acids in the pre-S region is linked to a carrier via active sites on the carrier. Still more preferred, the carrier is a lipid vesicle carrier. Even more preferred, the lipid vesicle carrier is stabilized by cross-linking.

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_	The carrier of the present invention includes a
2	lipid vesicle stabilized by cross-linking and having
3	covalently bonded active sites on the outer surface thereof.
4	The synthetic peptide is bonded via such active sites on the
5	carrier to the outer surface of the lipid vesicle. Such
6	active sites include COOH, -CHO, -NH2 and -SH. Such
7	stabilization by cross-linking is accomplished by a
8	
9	stabilizing agent such as an aldehyde having at least two
10	functional groups, such as a bifunctional aldehyde, for
11	example, glutaraldehyde. The carrier of the present
12	invention is chemically cross-linked with pendant functional
13	groups.
14	The present application also concerns diagnostic
15	methods. The present invention relates to processes for
16	detecting the presence of either pre-S gene coded -
17	hepatitis B antigens or antibodies in a serum.
18	Antibodies to the synthetic peptides disclosed
19	herein can be detected in samples by a process which
20	comprises:
21	<ul> <li>a) contacting the sample with a solid substrate</li> </ul>
22	coated with a non-labelled peptide containing an amino acid
23	chain corresponding to at least six consecutive amino acids
24	within the pre-S gene coded region of the envelope of HBV,
25	the peptide free of an amino acid sequence corresponding to
26	the naturally occurring envelope proteins of hepatitis B
27	virus, incubating and washing said contacted sample;
28	b) contacting the incubated washed product
29	obtained from step a above with a labelled peptide
30	

containing an amino acid chain corresponding to at least six consecutive amino acids within the pre-S gene coded region of the envelope of HBV, said peptide free of an amino acid sequence corresponding to the naturally occurring envelope protein of hepatitis B virus, incubating and washing the resultant mass; and

c) determining the extent of labelled peptide present in the resultant mass obtained by step b above.

Such a process is normally performed using a solid substrate which is substantially free of available protein binding sites. Such as by binding sites unbound by unlabelled peptide with a protein binding site occupier, e.g., albumin.

Another process for detecting such antibodies comprises:

- a) contacting the sample with a solid substrate coated with a non-labelled peptide containing an amino acid chain corresponding to at least six consecutive amino acids within the pre-S gene coded region of the envelope of HBV, the peptide free of an amino acid sequence corresponding to the naturally occurring envelope proteins of hepatitis B virus, incubating and washing said contacted sample;
- b) contacting the incubated washed product obtained from step a above with labelled antibody to human or animal immunoglobulin product by contact with an immunogen comprising a peptide corresponding to at least six consecutive amino acids within the pre-S gene coded region of the envelope of HBV, the peptide immunogen free of an amino acid sequence corresponding to the naturally occurring

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1	envelope prote	ins of hepatitis B virus,	incubating and
2	washing the co	ntacted sample, and	
3	c)	determining the extent of	f labelled antibody
4	present in the	resultant mass of step b	•

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HBV or HBsAg can be detected in a sample by a process which comprises:

- a) contacting a first portion of a composition containing an antibody produced by introducing into an animal or human an immunogen comprising a peptide corresponding to at least six consecutive amino acids within the pre-S gene coded region of the envelope of HBV, the peptide immunogen free of an amino acid sequence corresponding to the naturally occurring envelope proteins of hepatitis B virus, with a mixture of said sample and the immunogen which has been labelled, incubating and washing the first portion;
- b) contacting a second portion of the composition containing antibody with the same amount of the labelled immunogen in an antigen free control, incubating and washing the second portion;
- c) adding the same amount of Staphylococci
  bearing protein A to each of the compositions of steps a and
  b above, incubating both of the compositions, centrifuging
  each of the compositions and separating liquid from the
  solids therein;
- d) determining the extent of labelled immunogen in each of the resultant compositions from step c above, and
- e) comparing the relative amount of labelled immunogen in each such that if the activity of the resultant

composition containing the first portion is less than the activity for the resultant composition of the second portion, then the sample contains HBV or HBsAg.

 The synthetic immunogens can be used in general in both sandwich type immunoassays and competition type immunoassays, such as those immunoassays in which antigen in the sample competes with labelled immunogen for antibody.

These and other suitable immunoassay schemes for use in connection with the synthetic immunogens of this invention and antibodies thereto are disclosed in copending application Serial No. 426,309, filed September 29, 1982, entitled Labelled Peptides As Diagnostic Reagents, assigned to one of the assignees hereof, the disclosure of which is hereby incorporated herein by reference.

The present invention also concerns a diagnostic test kit for detecting hepatitis B virus in sera comprising

- a) antibodies to a peptide containing an amino acid chain corresponding to at least six consecutive amino acids within the pre-S gene coded region of the envelope of HBV, the peptide being free of an amino acid chain corresponding to the naturally occurring envelope proteins of hepatitis B virus, attached to a solid support,
- c) labelled antibodies to the peptide or to hepatitis B virus.

The kit can comprise a set of instructions for effecting an immunoassay wherein the effect of formation of an immune complex is revealed by said labelled antibody.

The present invention also concerns a diagnostic kit for detecting the presence of antibodies to pre-S gene

•	coded antigens of hepatitis B virus in a test sample
2	comprising
3	a) a given amount of a peptide containing
4	an amino acid chain corresponding to at least six
5	consecutive amino acids within the pre-S gene coded region
6	of the envelope of HBV, the peptide being free of an amino
7	acid chain corresponding to the naturally occurring envelope
8	proteins of hepatitis B virus. The petide is attached to a
9	solid support, e.g., a water insoluble solid support.
10	b) labelled antibodies, e.g., radiolabeled
11	or enzyme labelled, to human IgG and/or IgM.
12	The kit can comprise a set of instructions for
13	effecting an immunoassay, wherein the extent of formation of
14	an immune complex is revealed by said labelled antibodies.
15	In a particular aspect, the present invention
16 17	concerns a process for the detection of antigens coded for
18	the pre-S gene in sera of HBV infected humans and certain
19	animals, for example, chimpanzees, comprising the following
20	steps:
21	(a) coating a solid substrate with
22	antibodies to a peptide having an amino acid chain
23	corresponding to at least six consecutive amino acids within
24	the pre-S gene of HBV DNA, the peptide being free of an
25	amino acid sequence corresponding to the naturally occurring
26	envelope proteins of HBV,
27	(b) washing the coated substrate;
28	(c) contacting the washed coated substrate,
29	e.g., polystyrene beads; with a protein-containing solution;
30	(d) washing the substrate from step c:

1	(e) incubating the substrate from step d
2	with a sample suspected to contain HBV or HBsAg;
3	(f) washing the substrate from step e;
4	(g) adding radiolabeled or enzyme-labeled
5	antibody, the antibody being an antibody to the peptide or
6	HBsAg;
7	(h) incubating the substrate from step g;
8	(i) washing the substrate from step h; and
9	
10	counting in a gamma counter, or measuring its enzymatic
11	
12	activity.
13	The above process can be conducted using ELISA
14	techniques rather than RIA detection techniques.
15	In a particular embodiment, the present invention
16	also relates to a process for the detection of antibodies to
17	proteins coded for by the pre-S region of hepatitis B virus
18	DNA, comprising the following steps:
19	(a) adsorbing on a solid substrate
2 0	containing binding sites thereon, e.g., polystyrene beads, a
21	peptide having an amino acid sequence corresponding to at
22	least six consecutive amino acids within the pre-S gene
23	coded region of the HBV envelope, the peptide being free of
24	an amino acid sequence corresponding to the naturally
25	occurring envelope proteins of hepatitis B virus,
26	(b) contacting the substrate from step a
27	with a material to saturate the binding sites thereon,
28	(c) washing the substrate from step b,
29	(d) contacting the substrate from step c
30	with a specimen comprising human sera.

1	(e) incubating the resultant mass of cep d,
2	(f) washing the resultant mass of step e,
3	(g) adding radiolabeled antibodies to human
4	IgG or IgM to the resultant mass of step f to form a second
5	resultant mass,
6	(h) subjecting the second resultant mass of
7	step g to counting in a gamma counter,
8	(i) subjecting normal sera utilized as a
9	control to steps (a) to (h) and
10	(j) comparing the counts of steps h and i.
11	In the above process for the detection of
12	antibodies, ELISA techniques can be substituted for RIA
13	techniques.
14	The present invention also relates to a process
15	for predicting the outcome of hepatitis B infection which
16	
17	comprises carrying out an immunoassay on serum of a human to
18	detect the presence of an antibody to an antigen coded for
19	by the pre-S gene coded region of the envelope of hepatitis
20	B virus employing the above-described hepatitis B peptide
21	immunogen at regular intervals and evaluating the data.
22	The present invention further relates to a process
23	for determining if a human who has been vaccinated with a
24	vaccine against hepatitis B has become immune to hepatitis B
25	virus. Such process involves effecting a plurality of
26	immunoassays of serum from such human to determine if there
27	are antibodies in the serum to an antigen coded by the pre-S
28	gene coded region of the envelope of hepatitis B virus
29	employing the above-described hepatitis B peptide immunogen,
30	the immunoassays being performed on serum taken from the

human at different times.

The present invention further concerns a method for detecting the presence of hepatitis B virus infection comprising effecting quantitative immunoassays on a serum sample taken from a human to determine the amount of antibodies present therein which are antibodies to an antigen coded by the pre-S gene coded region of the envelope of the hepatitis B virus employing the above-described hepatitis B peptide immunogen and comparing the value with a known standard.

The present invention further concerns a method for detecting the presence of hepatitis B virus infection comprising effecting quantitative immunoassays on a serum sample taken from a human to determine the amount of antigens coded by the pre-S gene coded region of the envelope of the hepatitis B virus employing the abovedescribed antibodies to the hepatitis B peptide immunogen and comparing the value with a known standard.

The present invention also related to a process for raising antibodies which involves introducing into an animal the above-described hepatitis B peptide immunogen.

Still further, the present invention concerns a process for synthesizing His and Trp containing peptides
which includes the steps of

- a. linking a first amino acid containing an alpha-amino protecting group to a resin;
  - b. removal of the alpha-amino protecting group;
- 29 c. coupling a second amino acid containing an alpha-amino protecting group to the first amino acid;

1	d. repeating steps b and c by coupling further
2	alpha-protected amino acids to produce a desired peptide,
3	wherein at least one of the amino acids is His and wherein
4	at least one of said amino acids is Trp,
5	e. cleaving the peptide from the resin and
6	removing remaining protective groups to said first amino
7	acids;
8	<pre>f. substituting a His(ImDNP) for the His;</pre>
9	g. substituting a Trp(InFormyl) for the Trp;
10	h. removing the DNP prior to the cleavage and
11	the removing of protective groups, and
12	i. removing the Formyl during the cleavage and
13	the removing of protective groups.
14	
15	The present invention further concerns a
16	prophylatic method of protecting a patient against becoming
17	infected with hepatitis B comprising administering to such
18	patient, e.g., a human, an effective dosage of a vaccine as
19	described hereinabove
20	BRIEF DESCRIPTION OF THE DRAWINGS
21	Fig. 1 shows the results of submitting reduced
22	HBsAg disassociated into its constituent polypeptides to
23	SDS-polyacrylamide gel electrophoresis ("SDS-PAGE") in urea.
24	Panel a shows the separated proteins detected by a silver
25	stain and panel b is a Western blot with human antiserum to
26	hepatitis B.
27	Fig. 2 shows amino acid sequences of the
28	translational products of the pre-S gene region deduced from
29	sequences of HBV DNA. The sequences are presented in
30	one-letter amino acid code words (such code words are

defined in the Definitions herein). Sequences for five distinct HBV subtypes are presented. The 6th bottom line shows amino acid residues common to all five subtypes.

Fig. 3 shows a profile of relative hydrophilicity corresponding to the amino acid sequence of the pre-S gene product. Profiles for subtypes other than ayw are similar. The portion of the profile to the right from methionine 175 represents the S-gene translation product.

Fig. 4 shows two sets of bar graphs for mean antibody responses of rabbits immunized with free pre-S 120-145 (Fig. 4A) and with the same peptide linked to cross-linked liposomes containing L-tyrosine-azobenzene -p-arsonate (RAT) groups (Fig. 4B). Anti-HBs (antibodies to HBsAg), cross-hatched columns; anti-pre-S 120-145, diagonally hatched columns. Similar results to Fig. 4B were obtained with liposomes lacking RAT groups, except that responses after six weeks were lower. Columns corresponding to time = 0 represent sera before immunization.

Fig. 5 depicts radioimmunoassays with serial dilutions of a serum from a rabbit immunized with pre-S 120-145 linked to liposomes. Anti-HBs (antibodies to HBsAg), g; anti-pre-S 120-145, . Counts per minute (cpm) corresponding to distinct dilutions of the pre-immune serum were subtracted from cpm corresponding to dilutions of anti-pre-S 120-145; the difference was plotted. The endpoint titer of the serum (1/163,840) corresponds to its highest dilution at which the cpm were > 2.1 higher than those corresponding to the same dilution of the pre-immune serum.

Fig. 6 shows the reaction of anti-pre-S 120-145 1 with P33 and P36 in a Western blot (similar to Fig. 1). 2 Fig. 7 shows a graph depicting a diagnostic test 3 for hepatitis B antigens based on polystyrene beads coated 5 with anti-pre-S 120-145. Fig. 8 depicts a plot representing the compilation 6 of antibody responses of individual rabbits to conjugates of 7 S135-155 (amino acids 309 to 329 of the open reading frame 8 of the HBV env gene). The type of conjugates is indicated by 9 numbers defined in Table 1. Antibodies in sera obtained two 10 weeks after the third immunization were assayed using a 11 S135-155- beta-galactosidase conjugate and Pansorbin 12 (Neurath et al., 1982, supra). Their relative titer is given 13 in comparison with antibody levels induced by a S135-155-KLH 14 conjugate. Results of anti-HBs assays by RIA (AUSRIA test, 15 16 Abbott Laboratories, North Chicago, Illinois) are given in 17 international milliunits (mIU/ml; Neurath et al., 1982 18 supra). The line corresponds to the calculated linear 19 regression that best fits the set of all data concerning 20 rabbits with an anti-HBs response. The calculated 21 correlation coefficient (= 0.55) indicates a poor 22 correlation between anti-HBs and anti-S135-155 responses. 23 Fig. 9 shows four sets of bar graphs (Fig. 9A, 24 Fig. 9B. Fig. 9C and Fig. 9D) depicting examples of time 25 courses of antibody responses in rabbits immunized with 26 distinct S135-155-conjugates (indicated by numbers in each 27 panel and defined in Table 1). Fig. 9A corresponds to 28 conjugate No. 5; Fig. 9B corresponds to conjugage No. 11; 29 Fig. 9C corresponds to conjugage No. 12 and Fig. 9D

corresponds to conjugate No. 19. Anti-HBs (dashed columns) 2 and anti-S-135-155 (black columns) were assayed as described 3 for Fig. 8. Fig. 10 shows four plots (A, B, C and D) which 5 depict the kinetics of antibody responses to peptide 6 pre-S(120-145) (☑) and to pre-S protein within 7 approximately 22nm spherical HBsAG particles ( ) elicited 8 by unconjugated peptide pre-S(120-145) (plot A) and by the 9 same peptide linked to cross-linked, cysteine-activated 10 liposomes with attached RAT (L-tyrosine 11 azobenzene-p-arsenate) groups (plot B); and the effect of 12 carrier on anti-peptide antibody titers in sera of rabbits 13 immunized with 4 doses of peptides pre-S(120-145) (plot C) 14 and pre-S(12-32) (plot D) given 2 weeks apart. The carriers 15 for plots C and D were: (1) none; (2) keyhold lympet 16 hemocyanin (KLH); (3) alum; (4) and (5) cross-linked, 17 cysteine-activated liposomes with or without attached RAT 18 groups. Complete and incomplete Freund's adjuvant was used 19 in all cases except(3) . 20 Fig. 11 shows two plots for radioimmunoassays of 21 IgG antibodies in serial dilutions of rabbit antisera: to 22 pre-S(120-145) [ ]; to HBV particles and tubular forms of 23 THBSAg [O], devoid of antibodies to S-protein detectable by 24 RIA and to a fusion protein of chloramphenicol 25 acetyltransferase with the sequences of pre-S protein 26 lacking the 41 C-terminal amino acid residues (  $\Box$  ); and of 27 IgG ( $\Delta$ ) and IgM ( $\Delta$ ) antibodies in serum of a patient 28 recovered from hepatitis B. The latter serum was drawn 29 before antibodies to the S-protein were detectable. Immulon 30

1 2 Removable strips (Dynatech Lab ratories) were coated with 2 20  $\mu$ g/ml of either free peptide pre-S(120-145) or 3、 pre-S(12-32) and post-coated with gelatin (2.5 mg/ml in 0.1 M Tris, pH 8.8). The conditions for coating and the double 5 antibody RIA are described in A.R. Neurath, S.B.H. Kent, I 6 Strik, Science, 224, 392 (1984) and A.R. Neurath, S.B.H. 7 Kent, N. Strick, Proc. Natl. Acad. Sci USA, 79, 7871 (1982) 8 Fig. 12 shows a plot depicting the inhibition of 9 the reaction of anti-pre-S(120-145) IgG (antiserum diluted 10 1:100) with a pre-S(120-145)-B-galactosidase conjugate by; 11 free peptide pre-S(120-145) [ ]; by 20 nm spherical HBsAc 12 particles [A] and by HBV particles [D]. The latter two 13 preparations contained the same concentration of HBsAg 14 S-protein as determined by radioimmunoassay (AUSRIA, Abbot-15 Laboratories). 16 Fig. 13 depicts a plot of titers of anti-pre-17 S(120-145) antibodies versus days of surveillance and 18 indicates the development of IgM [●] and IgG [图] 19 antibodies to the pre-S gene coded protein of HBV during 20 acute hepatitis B. 21 Fig. 14 shows a plot for radioimmunoassays of 22 various preparations containing HBV-specific proteins on 23 polystyrene beads coated either with anti-pre-S(120-145) I 24 (o, •, □ ) or with IgG from a rabbit antiserum against HBV 25 particles and tubular forms of HBsAg ( $\triangle$ ,  $\triangle$ ). The tested

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antigens were: HBV particles and tubular forms  $(\bullet, \triangle)$ ;

approximately 20 nm spherical particles of HBsAg isolated

from plasma  $(o, \Delta)$ ; and the latter particles treated with

pepsin (1 mg/ml HBsAg, 50 µg/ml pepsin in 0.1 M glycine-UCL, pH 2.2, 2 hours at 37°C) (□).

Fig. 15 depicts a plot for radioimmunoassays of polymerized albumin-binding sites associated with HBsAg isolated from human plasma and containing pre-S gene coded sequences (•) or with HBsAg produced in yeast transfected with recombinant DNA containing the HBV DNA S-gene and thus lacking pre-S gene coded sequences (o).

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#### DETAILED DESCRIPTION OF THE INVENTION

Amino acid sequences deduced from sequences of the pre-S portion of the env genes corresponding to several HBV subtypes (see Fig. 2) have the following properties distinct from those of the S-protein: (i) high hydrophilicity and high percentage of charged residues (E. Schaeffer, J.J. Sninsky, Pro. Natl. Acad. Sci. USA, 81, 2902 (1984)); (ii) absence of cysteine residues; (iii) the highest subtype-dependent variability among HBV DNA gene products; and (iv) little homology with analogous sequences corresponding to nonhuman hepadnaviruses (F. Galibert, T.N. Chan, E. Mandart, J. Virol., 41, 51, (1982)). These properties suggest that the pre-S gene coded portion of the -HBV envelope is exposed on the surface of the virion, is a target for the host's immune response and is responsible for the host range of HBV (limited to humans and some primates). Synthetic peptides and antibodies against them, having predetermined specificity offer the opportunity to explore the biological rule of the pre-S protein moiety of the HBV envelope.

ı	Cleavage of disulfide bonds within HBsAg results
2	in:
3 *	(a) a substantial decrease of bindiny of
4	polyclonal antibodies (G.N. Vyas, K.R. Rao, A.B. Ibrahim,
5	Sience, 178, 1300, (1972); N. Sukeno, R. Shirachi, J.
6	Yamaguchi, N. Ishida, <u>J. Virol.</u> , 9, 182, (1972); G.R.
7	Dreesman, F.B. Hollinger, R.M. McCombs, J.L. Melnick, J.
8	Gen. Virol. 19, 129 (1973); and A.R. Neurath, N. Strick, J.
9	Med. Virol., 6, 309, (1980)) and of some monoclonal
10	antibodies (J. Pillot, M.M. Riottot, C. Geneste, L.
11	Phalente, R. Mangalo, Develop. Biol. Stand., in press
12	(1984)) elicited by intact HBsAg, and
13	(b) reduction of immunogenicity (Y. Sanchez, I.
14	Ionescu-Matiu, J.L. Melnick, G.R. Dreesman, J. Med. Virol.
15	11, 115, (1983)). However, some epitopes are resistant to
16	reduction of disulfide bonds (M. Imai, A. Gotoh, K.
17	Nishioka, S. Kurashina, Y. Miyakawa, M. Mayumi, J. Immunol.,
18	112, 416, (1974)). These epitopes are common to all
19	antigenic subtypes of HBV, but their localization on
20	envelope components of HBV has not been determined. The
21	present invention takes advantage of the localization of
22	disulfide-bond independent antigenic determinants on the
23	N-terminal portion (coded for by the pre-S gene of HEV DNA)
24	of the minor HBsAg proteins P33 and P36, and on other
25	regions of proteins coded for by the pre-S gene.
26	These determinants represent the dominant epitopes
27	on reduced and dissociated HBsAg reacting with human
28	anti-HBs. They are mimicked with high fidelity by pra-S
29	120-145 which elicits antibodies to HBsAg about 400 times
3.0	120-145 Which efficits antibodies to mostly described

more efficiently than a synthetic peptide analogue corresponding to the S-gene (A.R. Neurath, S.B.H. Kent, and N. Strick, <u>Proc. Natl. Acad. Sci. USA</u>, <u>79</u>, 7871 (1982)). No precedent exists for such high levels of virus-recognizing antibodies to a synthetic peptide analogue of an HBV protein. These antibodies could be used in a diagnostic test permitting the direct detection of the pre-S gene coded antigenic determinants in serum of HBV carriers.

The pre-S gene is the most divergent among all regions of hepadnavirus genomes (F. Galibert, T.N. Chen, E. Mandart, J. Virol., 41, 51 (1982)) (HBV is a member of the hepadnavirus family).

The hepatitis B vaccine of the present invention contains a peptide, either a synthetic peptide (peptide produced by assembling individual amino acids by chemical means or by expression vectors (DNA route)) or a peptide derived from natural sources, such peptide having an amino acid chain corresponding to at least six consecutive amino acids within the pre-S gene coded region of the surface antigen of hepatitis B virus. Such chain can be, for example, at least 10, 15, 20, or 26 amino acids long. A preferred peptide according to one embodiment of the present invention is an amino acid chain disposed between sequence position pre-S 120 and pre-S 174, and more preferably such chain includes the N-terminal methionine at sequence position pre-S 120. A preferred peptide is an amino acid chain corresponding to the chain between sequence position pre-5 120 and pre-S 145, i.e., pre-S (120-145).

1	Preferred Positions of the chain include the			
2	following: (1) The amino acid chain entirely between and			
3 •	including sequence position pre-S 1 and pre-S 11 for			
4	subtypes adw <sub>2</sub> and adr, (2) between and including sequence			
5	positions pre-S 10 and pre-S 40, (3) between and including			
6	sequence positions pre-S 15 and pre-S 120, (4) between and			
7	including sequence position pre-S 15 and pre-S 55, and (5)			
8	between and including sequence position pre-S 90 and pre-S			
9	120. A particularly preferred chain according to the			
10	present invention has 26 amino acids, includes the			
11	N-terminal methionine at sequence position pre-S 120 and is			
12	disposed between sequence position pre-S 120 and pre-S 174.			
13	Preferred peptides according to the present			
14	invention include the following:			
15	(1) pre-S(12-32), wherein the sequence is (see			
16	Fig. 2) MGTNLSVPNPLGFFPDHQLDP for subtype adw2;			
17	(2) pre-S(120-145), wherein the sequence is (see			
18	Fig. 2) MQWNSTAFHQTLQDPRVRGLYLPAGG for subtype adw2;			
19	(3) pre-S(32-53), wherein the sequence is (see			
20	Fig. 2) PAFGANSNNPDWDFNPVKDDWP for subtype adw2;			
21	(4) pre-S(117-134), wherein the sequence is (see			
22	Fig. 2) PQAMQWNSTAFHQTLQDP for subtype adw <sub>2</sub> ;			
23	(5) pre-S(94-117), wherein the sequence is (see			
24	Fig. 2) PASTNRQSGRQPTPISPPLRDSHP for subtype adw <sub>2</sub> ;			
25	(6) pre-S(153-171), wherein the sequence is (see			
26				
27	Fig. 2) PAPNIASHISSISARTGDP for subtype adw <sub>2</sub> ;			
28	(7) pre-S(1-21), wherein the sequence is (see			
29	Fig. 2) MGGWSSKPRKGMGTNLSVPNP for subtype adw2;			
30				

(8) pre-S(57-73), wherein the sequence is (see 1 2 Fig. 2) QVGVGAFGPRLTPPHGG for subtype adw; 3 (9) pre-S(1-11), 4 a. for adw2, wherein the sequence is (see 5 Fig. 2) MGGWSSKPRKG 6 b. for adr, wherein the sequence is (see 7 Fig. 2) MGGWSSKPRQG. Я Any analogs of the pre-S gene coded sequences of 9 the present invention involving amino acid deletions, amino 10 acid replacements, such as replacements by other amino 11 acids, or by isosteres (modified amino acids that bear close 12 structural and spatial similarity to protein amino acids), 13 amino acid additions, or isosteres additions can be 14 utilized, so long as the sequences elicit antibodies 15 recognizing the pre-S protein of HBV or hepatitis B surface 16 antigen. 17 In the formation of a peptide derived from natural 18 sources, a protein containing the required amino acid 19 sequence is subjected to selective proteolysis such as by 20 splitting the protein with chemical reagents or using 21 enzymes. Synthetic formation of the peptide requires 22 chemically synthesizing the required chain of amino acids. 23 In forming a synthetic vaccine according to the 24 present invention, it is preferred to insure that the amino 25 acid chain (peptide residue) corresponding to at least six 26 consecutive amino acids within the pre-S gene coded region 27 of hepatitis B virus has the steric configuration to be 28 recognized by antibody to hepatitis B virus. To this end, 29 the given chain of amino acids may have bonded thereto as 30

part of the amino acid chain, one or more additional amino acids on either, or both sides thereof. These additional amino acids can serve as auxiliary amino acids to enhance the stabilization of the amino acid chain so that it is readily recognized by antibody to hepatitis B virus. The additional amino acids can be the same amino acids in the same sequence as they occur in the natural protein, or other amino acids may be employed.

In one form of the invention, the peptide having a chain length of minimally six amino acids can be bounded on either side thereof with additional amino acids, e.g., three amino acids on either side of the residue, to form a longer chain of amino acids. The chain of amino acids may contain more than one amino acid sequence corresponding to at least six consecutive amino acids within the pre-S region of the surface antigen of hepatitis B virus.

would depend on the method of producing the sequence. If the sequence is made by assembling individual amino acids by chemical means, then the sequence length would generally not exceed 50 amino acids, and preferably would not exceed 40 amino acids. If the synthetic peptide is obtained from a DNA route, the chain length could be longer, for example, 100 or more amino acids. It is, however, normally shorter, and optimally considerably shorter than the natural pre-S protein. Thus, in the embodiment wherein the peptide has units of both the S region and pre-S region, its peptide portions corresponding to the S region is shorter than the natural S protein, e.g., no more than 100 amino acids,

preferably no more than 40 amino acids and usually less than

30 amino acids. In such cases, the peptide portion

corresponding to the pre-S region can be of a length

corresponding to the entire pre-S region, but generally is

less than the entire pre S region.

When the peptide contains no components corresponding to the amino acid sequence of the S region, it can contain amino acid sequences corresponding to the entire pre-S region, or shorter than the entire pre-S region.

Where, however, the amino acid sequence is part of a long chain, such as when there are more than one sequence of amino acids, the chain can contain residues of various moieties, for example, segments of polyamino acids or polysaccharides.

In addition to containing one or more different or the same sequences of amino acids corresponding to at least six consecutive amino acids within the pre-S region of hepatitis B virus, e.g., containing more than one sequence of amino acids corresponding to different epitopes (antigenic determinants) in the pre-S region of hepatitis B virus, the vaccine of the present invention can contain amino acid chains containing epitopes of different antigens or allergens so as to form a vaccine directed to hepatitis B virus and to one or more additional diseases, e.g., measles, influenza, smallpox, polio, diptheria, just to name a few. Such additional amino acid sequences can be of varying amino acid chain lengths.

A hepatitis B vaccine according to the present invention can include in addition to one or more amino acid

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        sequences corresponding to at least six consecutive amino
2
        acids within the pre-S region of the surface antigen of
3
      - hepatitis B virus, one or more amino acid sequences
4
        corresponding to consecutive amino acids within the S region
5
        of the surface antigen of hepatitis B virus, for example,
6
                                            145
                                                  146
                               143
                                     144
                   141
                         142
                                            Gly
                                                  Asn,
                         Pro
                               Thr
                                     Asp
                  Lys
7
                                or
8
                                    143
                                                    146
                                                         147
                                                              148
                                                                    149
                                          144
                                               145
                               142
                          141
        138
              139
                     140
9
                              Pro Thr
                                         Asp
                                              Gly
                                                    Asn Cys
                                                              Thr
                                                                    Cys
                          Lys
                     Thr
              Cys
        Cys
10
                   Other peptides corresponding to antigenic
11
        determinants of HBsAg (S region) and thus combinable in the
12
        same chain with one or more amino acids sequences
13
        corresponding to at least six amino acids in the pre-S
14
        region of HBsAg include the following:
15
        (1)
              Ser — Thr — Gly — Pro — Ser
16
                                         121
              117
17
                                                 122
18
                                                 Thr
19
                                               = Cys
                                          Cys
                                                     - Met - Thr
                                   Ser-
20
                                          137
                                                 124
                                                              - 1
                                                              Thr
                                   Pro
21
                                                               -
                                                              Ala
                                   Tyr
22
                                   Met - Ser - Thr - Gly - Gln
23
                                   Amino Acid Series
              Position
         (2)
24
                          Cys-Leu-Gly-Gln-Asn-Ser-Gln-Ser-Pro-Thr-
              48-81
25
                          Ser-Asn-His-Ser-Pro-Thr-Ser-Cys-Pro-Pro-
26
                          Thr-Cys-Pro-Gly-Thr-Arg-Trp-Met-Cys-Leu-
27
                          Arg-Arg-Phe-Ile
28
29
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1	(3,	2-16	Glu-Asn-Ile-Thr-Ser-Gly-Phe-Leu-Gly-Pro-
2			Leu-Leu-Val-Leu-Gln-Cys
3			·
4	(4)	22-35	Leu-Thr-Arg-Ile-Leu-Thr-Ile-Pro-Gln-Ser-
5			Leu-Asp-Ser-Trp-Cys
6			
7	(5)	38-52	Ser-Leu-Asn-Phe-Leu-Gly-Gly-Thr-Thr-Val-
8			Cys-Leu-Gly-Gln-Asn
9			
10	(6)	47-52	Val-Cys-Leu-Gly-Gln-Asn
11	(0)	47-52	Val-cys-bed dif din Ash
12			
13	(7 <b>)</b>	95-109	Leu-Val-Leu-Leu-Asp-Tyr-Gln-Gly-Met-Leu-
14			Pro-Val-Cys-Pro-Leu
15	(8)	104-109	Leu-Pro-Val-Cys-Pro-Leu
16			sequences of amino acids can be interconnected
17	with	h one anothe	er such as by cross-linking or by being bonded
18	dir	ectly there	to in the form of a branched chain, or the
19	res	pective sequ	mences can be bonded to a central "carrier".
20		There	e is realized by the present invention a
21	syn	thetic vacc	ine which is characterized by the absence of
22	nat	urally occu	ring envelope proteins of hepatitis B virus,
23	i.e	., the vacc	ine of the present invention is composed of
24	one	or more pe	ptide sequences corresponding to a limited
25	_ por	tion of the	hepatitis B virus envelope protein. The
26	vac	cine of the	present invention is also free of other
			in the virion. Vaccines can be synthesized
27			of biologically produced components, free of
28			ts whether they be active or inactive, free of
29			
30	ant	inodies, ir	ee of deoxyribonucleic acid (DNA), and are

therefore likely to be substantially free from undesirable side effects commonly found with other vaccines (e.g., unintentional infection with virus, allergic reactions, fevers, etc.).

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It should be understood that the vaccine of the present invention can be in admixture with other proteins and these proteins include the proteins of known antigens or allergens. Thus when it is stated herein that the vaccine is characterized by the absence of an amino acid sequence corresponding to the naturally occurring envelope proteins of the hepatitis B virus it is meant that notwithstanding the absence of such proteins, the composition functions as a vaccine, i.e., provides protective immunization by formation of antibodies.

The peptide of the present invention is such that it is capable of forming "neutralizing antibodies", i.e., antibodies that will protect patients against hepatitis B virus. Accordingly, the present invention is also directed to methods for protecting a patient against contracting hepatitis B.

The peptides and vaccines of the present invention can be used to improve immune response and to overcome non-responsiveness to certain known hepatitis B virus vaccines (e.g., containing no peptides corresponding to amino acid sequences in the pre-S region).

The peptides of the present invention can be utilized in conjunction with peptides containing amino acid chains corresponding to consecutive amino acids within the S ....gene coded region of HBsAg. Also, embodied by the present

invention is a peptide containing amino acids corresponding to consecutive amino acids spanning both the pre-S and S region, e.g., pre-S 160 to S 20.

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A carrier may be provided for the synthetic peptide of the invention. It should be understood, however, that a carrier may not be required to practice the present invention, i.e., a carrier may not be required to produce antibodies according to the present invention.

The "carrier" is simply a physiologically acceptable mass to which the synthetic peptide is attached and which is expected to enhance the immune response. A carrier can comprise simply a chain of amino acids or other moieties and to that end it is specifically contemplated to use as a carrier a dimer, oligomer, or higher molecular weight polymer of a sequence of amino acids defining a synthetic peptide of the invention. In other words, having determined the desired sequence of amino acids to form the synthetic peptide, these amino acids can be formed from naturally available materials or synthetically and can be polymerized to build up a chain of two or more repeating units so that repeating sequences serve both as "carrier" and synthetic peptide. Stated differently, an independent carrier may not be required. Alternatively, additional amino acids can be added to one or both ends of the amino acid chain that defines the synthetic peptide. It is preferred that alternative carriers comprise some substance, animal, vegetable or mineral, which is physiologically acceptable and functions to present the synthetic peptide so that it is recognized by the immune system of a host and stimulates a

satisfactory immunological response. Thus, a wide variety of carriers are contemplated, and these include materials which are inert, which have biological activity and/or promote an immunological response. For instance, proteins can be used as carriers. Examples of protein carriers include tetanus toxoid, keyhole lympet hemocyanin, etc.

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polysaccharides are also contemplated as carriers, and these include especially those of molecular weight 10,000 to 1,000,000, including, in particular, starches, dextran, agarose, ficoll or its carboxy methyl derivative and carboxy methyl cellulose.

polyamino acids are also contemplated for use as carriers, and these polyamino acids include, among others, polylysine, polyalanyl polylysine, polyglutamic acid, polyaspartic acid and poly (C2-C10) amino acids...

Organic polymers can be used as carriers, and these polymers include, for example, polymers and copolymers of amines, amides, olefins, vinyls, esters, acetals, polyamides, carbonates and ethers and the like. Generally speaking, the molecular weight of these polymers will vary dramatically. The polymers can have from two repeating units up to several thousand, e.g., two thousand repeating units. Of course, the number of repeating units will be consistent with the use of the vaccine in a host animal. Generally speaking, such polymers will have a lower molecular weight, say between 10,000 and 100,000 (the molecular weight being determined by ultracentrifugation).

Inorganic polymers can also be employed. These inorganic polymers can be inorganic polymers containing

organi. moieties. In particular, silicates and aluminum hydroxide can be used as carriers. It is preferred that the carrier be one which is an immunological adjuvant. In such cases, it is particularly contemplated that the adjuvant be muramyl dipeptide or its analogs.

The carrier can also be the residue of a crosslinking agent employed to interconnect a plurality of synthetic peptide containing chains. Crosslinking agents which have as their functional group an aldehyde (such as glutaraldehyde), carboxyl, amine, amido, imido or azidophenyl group. In particular, there is contemplated the use of butyraldehyde as a crosslinking agent, a divalent imido ester or a carbodiimide. Particularly contemplated divalent imido esters are those of the formula

$$R - 0 - C = NH_2^+$$
 $(CH_2)m$ 
 $R - 0 - C = NH_2^+$ 

wherein m is 1 to 13 and R is an alkyl group of 1 to 4 carbon atoms. Particularly contemplated carbodiimides for use as crosslinking agents include cyclohexylcarboxiimide, ethyldimethylaminopropyl carbodiimide, N-ethylmorpholino cyclohexyl carbodiimide and diisopropyl carbodiimide.

Chemical synthesis of peptides is described in the following publications: S.B.H. Kent, <u>Biomedical Polymers</u>, eds. Goldberg, E.P. and Nakajima, A. (Academic Press, New York), 213-242,(1980); A.R. Mitchell, S.B.H. Kent, M. Engelhard, and R.B. Merrifield, <u>J. Org. Chem.</u>, <u>43</u>, 2845-2852, (1978); J.P. Tam, T.-W. Wong, M. Riemen, F.-S. Tjoeng, and R.B. Merrifield, <u>Tet. Letters</u>, 4033-4036,

1 (1979); S. Mojsov, A.R. Mitchell, and R.B. Merrifield, J. 2 Org. Chem., 45, 555-560, (1980); J.P. Tam, R.D. DiMarchi and R.B. Merrifield, Tet. Letters, 2851-2854, (1981); and S.B.H. Kent, M. Riemen, M. Le Doux and R.B. Merrifield, Proceedings 5 of the IV International Symposium on Methods of Protein Segrence Analysis, (Brookhaven Press, Brookhaven, N.Y.), in 7 press, 1981. Chemical Synthesis: In the so-called "Merrifield 9 solid phase procedure" the appropriate sequence of L-amino 10 acids is built up from the carboxyl terminal amino acid to 11 the amino terminal amino acid. Starting with the appropriate 12 carboxyl terminal amino acid attached to a polystyrene (or 13 other appropriate) resin via chemical linkage to a 14 chloromethyl group, benzhydrylamine group, or other reactive 15 group of the resin, amino acids are added one by one using 16 the following procedure. The peptide-resin is: 17 (a) washed with methylene 18 chloride; 19 (b) neutralized by mixing for 10 minutes at room 20 temperature with 5% (v/v) diisopropyl-21 ethylamine (or other hindered base) in 22 methylene chloride; 23 (c) washed with methylene chloride; 24 (d) an amount of amino acid equal to six times the 25 molar amount of the growing peptide chain is 26 activated by combining it with one-half as 27 many moles of a carbodiimide (e.g., 28 dicyclohexylcarbodiimide, or diisopropyl-29 carbodiimide) for ten minutes at 0°C, to 30

1		form the symmetric anhyc ide of the amino
2		acid. The amino acid used should be
3		provided originally as the N-alpha-tert.butyl-
4		oxycarbonyl derivative, with side chains
5		protected with benzyl esters (e.g. aspartic or
6		glutamic acids), benzyl ethers (e.g., serine,
7		threonine, cysteine or tyrosine),
8		benzyloxycarbonyl groups (e.g., lysine) or other
9		protecting groups commonly used in peptide
10		synthesis.
11	(e)	the activated amino acid is reacted with
12		the peptide-resin for two hours at
13		room temperature, resulting in addition
14		of the new amino acid to the end of the
15		
16		growing peptide chain.
17	(f)	the peptide-resin is washed with methylene
18		chloride;
19	(g)	the N-alpha-(tert. butyloxycarbonyl) group is
20		removed from the most recently added
21		amino acid by reacting with 30 to 65%, preferably
22		50% (v/v) trifluoroacetic acid in methylene
23		chloride for 10 to 30 minutes at room
24		temperature;
25	(h)	the peptide-resin is washed with methylene
26		chloride;
27	(i)	steps (a) through (h) are repeated until the
28		required peptide sequence has been
29		constructed.
30	The pepti	de is then removed from the resin and

simultaneously the side-chain protecting groups are removed, by reaction with anhydrous hydrofluoric acid containing 10% v/v of anisole or other suitable (aromatic) scavenger.

Subsequently, the peptide can be purified by gel filtration, ion exchange, high pressure liquid chromatography, or other suitable means.

In some cases, chemical synthesis can be carried out without the solid phase resin, in which case the synthetic reactions are performed entirely in solution. The reactions are similar and well known in the art, and the final product is essentially identical.

Isolation from natural sources: If sufficient quantities of the whole protein antigen are available, a limited portion of the molecule, bearing the desired sequence of amino acids may be excised by any of the following procedures:

- (a) Digestion of the protein by proteolytic enzymes, especially those enzymes whose substrate specificity results in cleavage of the protein at sites immediately adjacent to the desired sequence of amino acids;
- 24\_
- (b) Cleavage of the protein by chemical means. Particular bonds between amino acids can be cleaved by reaction with specific reagents. Examples include: bonds involving methionine are cleaved by cyanogen bromide;

asparaginyl-glycine bonds are cleaved by

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1	hydroxylamine;
2	(c) A combination of proteolytic and chemical
3	cleavages.
4	It should also be possible to clone a small
5	portion of the DNA, either from natural sources or prepared
6	by synthetic procedures, or by methods involving a
7	combination thereof, that codes for the desired sequence of
8	amino acids, resulting in the production of the peptide by
9	bacteria, or other cells.
10	Analogously, one can form chains containing a
11	plurality of amino acid sequences by the following
12	technique: An aqueous solution of a peptide or peptides is
13	mixed with a water-soluble carbodiimide (e.g., ethyl-
14	dimethyl-aminopropylcarbodiimide). This results in
15	polymerization of the peptide(s); depending on the use of
16	the side chain blocking groups mentioned above, either
17	straight chain or branched polymers of the peptide can be
18	made.
19	If desired the synthetic peptide of the present
20	invention can have bonded thereto a chain of any of the
22	following moieties: polypeptide, polyamino acid, poly-
23	saccharide, polyamide or polyacrylamide which can serve as a
_	stabilizing chain or as a bridge between amino acids of the
25	individual chains. Such chains are available commercially
26	or, in the case of polyamino acids, are formed by a process
27	which comprises: mixing a solution of the desired amino acid
28	sequence with a solution of the N-carboxylanhydride of the
-	

amino acid and allowing a base-catalyzed polymerization to

occur, which is initiated by the amine groups of the peptide.

Although a carrier may not be required, if a carrier is employed the deposition of a chain or chains on a "carrier" can be effected as follows:

- 1. Protein Carrier: The protein and the synthetic peptide are dissolved together in water or other suitable solvent, and covalently linked via amide bonds formed through the action of a carbodiimide. The resulting product may contain one or more copies of the peptide per protein monomer. Alternatively, the reduced peptide may be added to a carrier containing sulfhydryl groups to form disulfide bonds. Yet another method involves the addition of reduced peptide to protein carriers containing maleimidyl groups to form a covalent linkage by a Michael addition, or any other covalent attachment means.
- 2. Polysaccharide Carriers: Oligosaccharide carriers should have molecular weights in the range 1,000 to 1,000,000. In order to covalently link these to synthetic peptides, suitable functional groups must first be attached to them. Carboxyl groups may be introduced by reacting with iodoacetic acid to yield carboxymethylated polysaccharides, or by reacting with carbonyldiimidazole to yield activated carbonyl esters. Carboxymethyl polysaccharides are coupled to the peptide by a carbodimide reaction, while the activated carbonyl esters react spontaneously with peptides. Multiple copies of the synthetic peptide should be attached to each oligosaccharide unit.

3. Polyamino Acid Carriers: These carriers should have molecular weights in the range 1,000 to 1,000,000. Polylysine and polyornithine have primary amino groups on their side chains; polyaspartic acid and polyglutamic acid have carboxyl groups. Peptides may be coupled to these via amide bonds using the carbodimide reaction. Another carrier that provides amino groups for coupling is polylysine to which polyalanine can be attached to the side chains of the lysine residues. The synthetic peptide may be attached to the ends of polyalanine chains, also by a carbodiimide reaction. Multiple copies of the synthetic peptide should be attached to each oligopeptide unit.

The novel carrier of the present invention includes a lipid vesicle having active sites on the outer surface thereof. Such active sites include -COOH, -CHO, -NH2 and -SH. The lipid carrier can be stabilized by cross-linking by a stabilizing agent such as an aldehyde having at least two functional groups, such as a bifunctional aldehyde, e.g., glutaraldehyde.

The bonding of the peptide to the lipid vesicle carrier occurs at the active sites on the lipid vesicle on the exterior surface of the carrier. Without wishing to be bound by any theory of operability, it is believed that such bonding is at least covalent bonding.

It is possible to bind a peptide to two active sites on the outer surface of the lipid vesicle. For example, a -NH<sub>2</sub> group at one end of a peptide can bind with a -COOH active site on the outer surface of the lipid

vesicle. The other end of the peptide can then bind what another active site on the lipid vesicle, for example, a coon group on the other end of the peptide can bind with a -NH2 active site on the lipid vesicle.

The preferred carrier to support the synthetic peptides of the present invention is a lipid vesicle. Lipid vesicles can be formed by sonicating a lipid in an aqueous medium, by resuspension of dried lipid layers in a buffer or by dialysis of lipids dissolved in an organic solvent against a buffer of choice. The latter procedure is preferred. Lipid vesicles consist of spheres of lipid bilayers that enclose part of the aqueous medium.

Lipid vesicle (non-protein) carriers according to the present invention can be produced in a variety of ways. The preferred method to produce such carriers would be to treat a lipid vesicle containing aminoalkanes and diaminoalkanes having 10 to 18 carbon atoms, for example stearylamine, cetylamine and myrististylamine with a polyaldehyde, such as a dialdehyde, for example, butanedial (succinaldehyde), pentanedial (glutaraldehyde), hexanedial (adipoldehyde), heptanedial (pimelicaldehyde) and octanedial (suberaldehyde). Alternatively, a liposome containing aminoalkenes and diaminoalkenes having 10 to 18 carbon atoms, for example, oleylamine, can be treated with the aforementioned polyaldehydes. The lipid vesicle carrier thus formed has active aldehyde groups on the surface thereof allowing the direct linking of peptides via their N-terminal or lysine groups.

Peptides linked to lipid vesicle carriers

according to the present invention can also be prepared by treating an amino containing lipid vesicle as described above with a peptide activated by carbodiimide, for example, N-ethyl-N' (dimethylami..opropyl) carbodiimide.

Alternatively a carbodiimide activated peptide is linked to polyaldehyde, e.g., dialdehyde, treated lipid vesicles which have been further derivatized by reaction with a water-soluble diaminoalkane, e.g., ethylene diamine and propylene diamine.

Still further, lipid vesicles containing fatty acids (saturated and unsaturated) having 12 to 18 carbon atoms, e.g., stearic acid, oleic acid, palmitic acid and myristic acid, are activated with carbodiimide. Thereafter, the activated lipid vesicle is reacted with a peptide.

Another approach to form a carrier according to the present invention involves using a fatty acid aldehyde as a component of the lipid vesicle and treating such lipid vesicle as described for glutaraldehyde treated lipid vesicles. Such lipid vesicle reacts directly with amino groups of peptides.

In a preferred embodiment of a carrier according -to the present invention, the aforementioned lipid vesicle carrier formed by treating a amino or diaminoalkane (or amino or diaminoalkane) having 10 to 18 carbon atoms with a polyaldehyde is further reacted with cysteine (L-or D- or LD- cysteine). These lipid vesicles are then reacted with a peptide having -SH groups, i.e., cysteine containing

R

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pel ides. The link between the lipid vesicle and the 1 peptide is mediated by a disulfide bond. 2 Alternatively, a fatty acid mercaptan is used as a 3 component of the lipid vesicle, for example, 4 octadecanethiol. A cysteine containing peptide is directly 5 linked to such lipid vesicle. 6 Another approach to form carriers according to the 7 present invention involves the preparation of the above 8 described fatty acid mercaptan containing lipid vesicles 9 which are further reacted with a dimaleimide, for example, 10 para or ortho N-N'-phenylenedimaleimide. Such lipid vesicle 11 is then reacted with a cysteine containing peptide. 12 Alternatively, the link between the appropriate 13 lipid vesicle and the appropriate peptide can be 14 accomplished by commercially available cross-linking 15 reagents such as dimethyl adipimidate; dimethyl 16 3,3'-dithiobis-propionimidate; 2-iminothiolane; 17 di-succinimidyl suberate; bis 2-(succinimidooxy 18 carbonyloxy) -ethyl] sulfone; disuccinimidyl tartarate; 19 dithiobis (succinimidyl propionate); ethylene glycol 20 bis(succinimidyl succinate); N-5-azido-2-nitrobenzoyloxy-21 succinimide; p-azidophenacyl bromide; p-azido-phenylglyoxal 22 4-fluoro-3-nitrophenyl azide; N-hydroxysuccinimidyl-4-azide 23 benzoate; N-hydroxysuccinimidyl-4-azidosalicylic acid; m-24 maleimidobenzoyl N-hydroxy succinimide ester; methyl-4-25 azidobenzoimidate; p-nitrophenyl 2-diazo-3,3,3-trifluoro-26 proprionate; N-succinimidyl-6 (4'-azido-2'-nitrophenylamino 27 nexanoate; succinimidyl 4-(N-maleimidomethyl) cyclohexane-28 29 1-carboxylate; succinimidyl 4-(p-maleimidomethyl) butyrate; 30

1 N-(4-azidophenylthio)phthalimide; ethyl 4-azidophenyl 1, 2 4-dithiobutyrimidate: N-succinimidyl (4-azidophenyldithio) 3 propionate; 1-5-difluoro-2, 4-dinitrobenzene; 4 4,4'-difluoro-3,3'-dinitrodiphenyl-sulfone; 5 4,4'-diisothiocyano-2,2'-disulfonic acid stilbene; 6 p-phenylenediisothiocyanate; 4,4'-dithiobisphenylazide; 7 erythritolbiscarbonate; N-succinimidyl 3-(2-pyridyldithiol) 8 propionate; dimethyl pimelimidate and dimethyl suberimidate. 9 The lipid vesicles according to the present 10 invention act not only as carriers, but also as adjuvants. 11 The lipid vesicle synthetic carriers of the 12 present invention can be utilized to bind synthetic peptide 13 analogues (eliciting protective antibodies) of various 14 viral, bacterial, allergen and parasitic proteins of man and 15 animals, besides synthetic peptide analogues of hepatitis B 16 surface antigen, and especially the novel synthetic peptide 17 analogue of hepatitis B surface antigen containing amino 18 acid sequences corresponding to amino acid sequences in 19 pre-S gene coded region of the HBV. 20 Accordingly, the lipid vesicle synthetic carriers 21 of the present invention can be used to bind with synthetic 22 peptide analogues of the following viruses: influenza 23 hemagglutinin (A/memphis/102/72 strain, A/Eng 1878/69 24 strain, A/NT/60/68/29c strain, and A/Qu/7/70 strain), fowl 25 plague virus hemagglutinin, vaccinia, polio, rubella, 26 cytomegalovirus, small pox, herpes simplex types I and II, 27 yellow fever, Infectious ectromelia virus, Cowpox virus, 28 Infectious bovine rhinotracheitis virus, Equine rhino-29 pneumonitis (equine abortion) virus, Malignant catarrh virus 30

of cattle, Feline rhinotracheitis virus, Camine herpes 1 virus, Epstein-Barr virus (associated with infectious 2 mononucleosis and Burkitt lymphoma), Marek's disease virus, Sheep pulmonary adenomatosis (Jaagziekte) virus, Cytomegaloviruses, Adenovirus group, Human papilloma virus, 5 Feline panleucopaenia virus, Mink enteritis virus, African 6 horse sickness virus (9 serotypes), Blue tongue virus (12 7 serotypes), Infectious pancreatic necrosis virus of trout, Fowl sarcoma virus (various strains), Avian leukosis virus 9 (visceral, erythroblastic and myeloblastic), Osteopetrosis 10 virus, Newcastle disease virus, Parainfluenza virus 1, 11 Parainfluenza virus 2, Parainfluenza virus 3, Parainfluenza 12 4, Mumps virus, Turkey virus, CANADA/58, Canine distemper 13 virus, Measles virus, Respiratory syncytial virus, 14 Myxovirus, Type A viruses such as Human influenza viruses, 15 e.g., Ao/PR8/34, A1/CAM/46, and A2/Singapore/1/57; Fowl 16 17 plaque virus; Type B influenza viruses, e.g., B/Lee/40; 18 Rabies virus; Eastern equinine encephalitis virus; 19 Venezuelan equine encephalitis virus; Western equine 20 encephalitis virus; Yellow fever virus, Dengue type l virus 21 (=type 6), Dengue type 2 virus (=type 5); Dengue type 3 22 virus; Dengue type 4 virus; Japanese encephalitis virus, 23 Kyasanur Forest virus; Louping ill virus; Murray Valley 24 encephalitis virus; Omsk haemorrhagic fever virus (types I 25 and II); St. Louis encephalitis virus; Human rhinoviruses, 26 Foot-and-mouth disease virus; Poliovirus type 1; Enterovirus 27 Polio 2; Enterovirus Polio 3; Avian infectious bronchitis 28 virus; Human respiratory virus; Transmissible 29 gastro-enteritis virus of swine; Lymphocytic 30

choriomeningitis virus; Lassa virus; Machupo virus; Pichinde virus; Tacaribe virus; Papillomavirus; Simian virus; Sindbis virus, and the like.

The lipid vesicle synthetic carriers of the present invention can be used to bind synthetic peptide analogues of bacteria, for example, leprosy, tuberculosis, syphilis and gonorrhea.

The lipid vesicle synthetic carriers of the present invention can also be used to bind synthetic peptide analogues of the following parasites: organisms carrying malaria (P. Falciparum, P. Ovace, etc.), Schistosomiasis, Onchocerca Volvulus and other filiarial parasites, Trypanosomes, Leishmania, Chagas disease, amoebiasis, hookworm, and the like.

The lipid vesicle carriers of the present invention can be used to bind the novel peptides of the present invention corresponding to amino acid sequences in the pre-S region of HBsAg. The lipid vesicle carriers of the present invention can also be used to bind amino acid sequences in the S region, as well as other amino acid sequences for other virus, etc.

Amino acid sequences (corresponding to amino acids in the S region) which contains an antigenic determinant for hepatitis B surface antigen can be linked to the lipid vesicle carrier of the present invention. T.P. Hopp, "A Synthetic Peptide with Hepatitis B Surface Antigen Reactivity", Mol. Imm., 18, 9, 869-872, 1981, propose the following sequence corresponding to the S region of HBsAg: 138 139 140 141 142 143 144 145 146 147 148 149

```
Cys Cys Thr Lys Pro Thr Asp Gly Asn Cys Thr Cys
1
                  Other peptides mimicking the antigenic determinant
2
        of HBsAg (S region) include the following:
3
4
        (1)
5
             Peptide 1
6
7
                            Thr
 8
                          = Cys - Met - Thr
 9
                                        Thr
10
              Pro
                                         Ala
11
              Tyr
              Met - Ser - Thr - Gly-
12
 13
         Peptide 2 contains 5 additional amino acid residues:
 14
              Ser - Thr - Gly - Pro - Ser - X,
 15
               117
         G.R. Dreesman, Y. Sanchez, I. Ionescu-Matiu, J. T. Sparrow,
 16
          H. R. Six, D.L. Peterson, F.B. Hollinger and J.L. Melnick,
 17
          "Antibody to Hepatitis B Surface Antigen After A Single
 18
          Inoculation of Uncoupled Synthetic HBsAg Peptides", Nature,
 19
          295, 158-160, 1982; and (2) the following peptides:
  20
  21
  22
                               SEQUENCE
          POSITION
                          Cys-Leu-Gly-Gln-Asn-Ser-Gln-Ser-Pro-Thr-Ser-
  23
            48-81
                          Asn-His-Ser-Pro-Thr-Ser-Cys-Pro-Pro-Thr-Cys-
  24
                          Pro-Gly-Tyr-Arg-Trp-Met-Cys-Leu-Arg-Arg-Phe-
  25
  26
                          Ile
                          Glu-Asn-Ile-Thr-Ser-Gly-Phe-Leu-Gly-Pro-Leu-
   27
             2-16
   28
                           Leu-Val-Leu-Gln-Cys
   29
   30
```

1	22-35	Le: Thr-Arg-Ile-Leu-Phr-Ile-Pro-Gln-Ser-Leu-	
2	•	Asp-Ser-Trp-Cys	
3			
4	38-52	Ser-Leu-Asn-Phe-Leu-Gly-Gly-Thr-Thr-Val-Cys-	
5		Leu-Gly-Gln-Asn	
6			
7	47-52	Val-Cys-Leu-Gly-Gln-Asn	
8		*	
9	95-109	Leu-Val-Leu-Leu-Asp-Tyr-Gln-Gly-Met-Leu-Pro-	
10	30 103	Val-Cys-Pro-Leu	
11		<b>142 6/4 12</b> 6 868	
12	104-109	Leu-Pro-Val-Cys-Pro-Leu	
13		ti-influenza Response Achieved by Immunization	
14	•		
15	<del>-</del>	tic Conjugate", Proc. Natl. Acad. Sci. USA, 79	_,
16	•	. The peptide corresponds to the sequence	
17	serine-91 to	leucine-108 of the amino acid chain of the	
18	virus.		
19	A p	eptide containing an amino acid sequence	
20	mimicking the	antigenic determinant of polyoma virus medium	n
21	size tumor an	tigen is Lys-Arg-Ser-Ars-His-Phe, G. Walter,	
22	M.A. Hutchins	on, T. Hunter and W. Eckhart, "Purification of	Ē
23	Polyoma Virus	Medium-Size Tumor Antigen by Immunoaffinity	
24	Chromatograph	y", Proc. Natl. Acad. Sci USA, 79, 4025-4029,	
25	1982.		
26	A p	eptide containing an amino acid sequence	
27	mimicking the	antigenic determinant of poliovirus replicase	e
28	antigen is as	follows:	
29	Tyr	-Ser-Thr-Leu Tyr-Arg-Arg-Trp-Leu-Asp-Ser-Phe	
30	450	461,	

1	M. H. Baron and D. Baltimore, "Antibodies Against a
2	Synthetic Peptide of the Poliovirus Replicase Protein:
3	Reaction with Native, Virus-Encoded Proteins and Inhibition
4	of Virus-Specific Polymerase Activities In Vitro". Jour.
5	Virology, 43, 3969-3978, 1982.
6	Peptides containing an amino acid sequence
7	mimicking the antigenic determinant of simian virus 40 large
8	tumor antigen are as follows:
9	Met-Asp-Lys-Val-Leu-Asn-Arg and
10	Lys-Pro-Pro-Thr-Pro-Pro-Pro-Glu-Pro-Glu-Thr,
11	G. Walter, K.H. Scheidtmann, A. Carbone, A.P. Laudano and
12	R.A. Lerner, N. Green, H. Alexander, FT. Liu, J.G.
13	Sutcliffe and T.M. Shinnick, "Chemically Synthesized
14	Peptides Predicted From the Nucleotide Sequence of the
15	Hepatitis B Virus Genome Elicit Antibodies Reactive With the
16	Native Envelope Protein of Dane Particles", Proc. Natl.
17	Acad. Sci. USA, 78, 6, 3403-3407, 1981.
18	A peptide containing an amino acid sequence
19 20	mimicking the antigenic determinant of retrovirus R antigen
21	is as follows:
22	Leu-Thr-Gln-Gln-Phe-His-Gln-Leu-Lys-Pro
23	Ile-Glu-Cys-Glu-Pro,
24	J.G. Sutcliffe, T.M. Shinnick, N. Green, FT. Liu, H.L.
25	Niman and R.A. Lerner, "Chemical Synthesis of A Polypeptide
26	Predicted From Nucleotide Sequence Allows Detection Of A New
27	Retroviral Gene Product, Nature, 287, 1980.
28	A peptide containing an amino acid sequence
29	mimicking the antigenic determinant of avian sarcoma virus
30	antigen is as follows:

```
1
                   Glu-Asp-Asn-Glu-Tyr-Thr-Ala-Arg-Gln-Gly,
 2
        `T.W. Wong and Alan R. Goldberg, "Synthetic Peptide Fragment
 3
        Of src Gene Product Inhibits the src Protein Kinase and
        Cross reacts Immunologically With Avian onc Kinases and
        Cellular Phosphoproteins", Proc. Natl. Acad. USA, 78, 12,
 6
        7412-7416, 1981.
 7
                   Peptides containing an amino acid sequence
 8
        mimicking the antigenic determinant of foot-and-mouth
 9
        disease virus antigen are as follows:
10
             141
             Val
                       Asn Leu Arg Gly Asp Leu Gly
11
12
                                                           160
                       Gly Lys
                                 Val Ala Arg
                                                 Thr
                                                           Pro
13
                             and
14
             201
15
                  Lys Gln Lys Ile Val Ala Pro Val
             His
                                                          Lys Gln
16
             Thr
                  Leu,
17
        J.L. Bittle, R.A. Houghten, H. Alexander, T.M. Shinnick,
18
        J.G. Sutcliffe, R.A. Lerner, D.J. Rowlands and F. Brown,
19
        "Protection Against Foot-And-Mouth Disease By Immunization
20
        With A Chemically Synthesized Peptide Predicted From the
21
        Viral Nucleotide Sequence", Nature, 298, 30-33, 1982.
22
                  A peptide containing an amino acid sequence
23
        mimicking the antigenic determinant of hemagglutinin X-31
24
        (H3N2) influenza virus antigen is as follows:
25
                          125
                  Glu-Gly-Phe-Thr-Trp-Thr-Gly-
26
                                       135
27
                  Val-Thr-Gln-Asn-Gly-Gly-Ser-
28
                  Asp Ala-Cys-Lys-Arg-Gly-Pro-
29
                      145
30
                  Gly-Ser-Gly-Phe-Phe-Ser-Arg-
```

•	151
2	Leu,
3	D.C. Jackson, J.M. Murray, D.O. White, C.N. Fagan and G.W.
4	Tregear, "Antigenic Activity of a Synthetic Peptide
5	Comprising the 'Loop' Region of Influenza Virus
6	Hemagglutinin", <u>Virology</u> , 120, 273-276, 1982.
7	A peptide containing an amino acid sequence
8	mimicking the antigenic determinant of hemagglutinin of type
9	A H3N2 influenza virus antigen was synthesized by G.M.
	Muller, M. Shapira and R.F. Doolittle, "Antibodies Specific
10	for the Carboxy- And Amino- Terminal Regions of Simian Virus
11	40 Large Tumor Antigen", Proc. Natl. Acad. Sci USA, 77, 9,
12	5179-5200, 1980.
13	A peptide containing an amino acid sequence
14	mimicking the antigenic determinant of influenza virus
15	strain 3QB antigen is Ile Val Asx Thr Ser Glx Pro1
16	Gly <sub>3</sub> Ala <sub>1</sub> Leu <sub>1</sub> Lys <sub>1</sub> , A. Aitken and C. Hannoun, "Purification
17	of Haemagglutinin and Neuraminidase from Influenza Virus
18	
19	Strain 3QB and Isolation of a Peptide From an Antigenic
20	Region of Haemagluttinin", Eur. J. Biochem, 107, 51-56,
21	1980.
22	Peptides containing an amino acid sequence
23	mimicking the antigenic determinant of diptheria antigen are
24	given as follows:
25	Natural DT Loop
26	-Cys-Ala-Gly-Asn-Arg-Val-Arg-Arg-Ser-Val- 186 190 195
27	Gly-Ser-Ser-Leu-Lys-Cys-
28	201
29	Synthetic Peptide
30	Tetradecapeptide Gly(188)Cys-(201)

1	Hexadecapeptide Cys(186)Cys-(201)
2	Octadecapeptide Ala-Ala-Cys(186)Cys-(201)
3	F. Audibert, M. Jolivet, L. Chedid, R. Arnon and M. Sela,
4	"Successful Immunization With a Totally Synthetic Diphtheria
5	Vaccine", Proc. Natl. Acad. Sci. USA, 79, 5042-5046, 1982.
6	A peptide containing an amino acid sequence
7	mimicking the antigenic determinant of Streptococcus
8	pyogenes M antigen is as follows:
9	5
10	Asn-Phe-Ser-Thr-Ala-Asp-Ser-Ala-Lys
11	10 15 Ile-Lys-Thr-Leu-Glu-Ala-Glu-Lys-Ala-Ala-
12	20 25
13	Leu-Ala-Ala-Arg-Lys-Ala-Asp-Leu-Glu-Lys-
14	30 - 35 Ala-Leu-Glu-Gly-Ala-Met
15	· · · · · · · · · · · · · · · · · · ·
16	E.H. Beachey, J.M. Seyer, D.B. Dale, W.A. Simpson and A.H.
17	Kang, "Type-Specific Protective Immunity Evoked by Synthetic
18	Peptide of Streptococcus Pyogenes M Protein", Nature, 292,
19	457-459, 1981.
20	The lipid vesicle carrier of the present invention
21	can thus be utilized to bind with any amino acid sequence
22	which includes the antigenic determinant for a specific
2.3	antigen.
24	The lipid vesicle carriers of the present
25	invention can also be used to bind with enzymes.
26	The present invention is also directed to
27	diagnostic tests for direct detection of HBV antigens and
28	HBV antibodies.
29	In order to detect HBV antigens containing
30	proteins coded for by the pre-S gene in sera of HBV-infected

1	animals such as humans, radioimmunoassay (RIA, Of
2	enzyme-linked immunosorbent assay (ELISA) can be employed.
3	. One test for detecting HBV antigens according to
4	the present invention is as follows:
5	(1) a solid substrate containing binding sites
6	thereon, e.g., polystyrene beads, is coated with antibodies
7	to a peptide having an amino acid chain corresponding to at
8	least six amino acids within the pre-S gene coded region of
9	the envelope of HBV, the peptide free of an amino acid
10	sequence corresponding to the naturally occuring proteins of
11	HBV;
12	(2) the coated beads are then washed with, for
13	example, tris buffered saline, to remove excess antibody;
14	(3) the beads are then contacted with a protein-
15	containing solution, such as bovine serum albumin (BSA) or
16	gelatin to saturate protein binding sites on the beads (to
17	prevent or reduce non-specific binding) - a convenient
18	concentration of such protein-containing solution can be
19	employed such as 1 mg/ml to 50 mg/ml;
20	(4) beads are then washed to remove excess BSA or
22	gelatin;
23	(5) the beads are then incubated with samples
24	suspected to contain HBV or HBsAg (normal sera is utilized
25	as a control);
26	(6) the beads are then washed with a solution,
27	e.g., tris buffered saline solution, and mixed with a
28	radiolabeled antibody, e.g., I labeled antibody (antibody
29	to either the peptide or to HBsAg);
30	(7) the beads are then incubated;

1 the beads are then washed and counted in a 2 gamma counter. 3 If the specimens have counts at least 2.1 times higher than counts of the control, then the specimens are 5 positive. 6 The pre-S gene coded peptides according to the 7 present invention can be employed as a diagnostic tool to 8 detect antibodies to the pre-S region of HBV in a given 9 sample. The pre-S gene coded peptide, for example, pre-S 10 (120-145), pre-S (12-32), pre-S (32-53), or pre-S (117-134), 11 pre-S(1-21), pre-S(94-117), pre-S(153-171), pre-S(32-53) and 12 pre-S(57-73), is adsorbed on a solid substrate, containing 13 binding sites thereon for example, polystyrene beads. The 14 substrate is thereafter contacted with a substance (protein 15 containing solution), for example, gelatin BSA or powdered 16 milk, to saturate the binding sites thereon. Thereafter, 17 the substrate is washed with a buffered solution and 18 thereafter the buffer is removed. A specimen, e.g., human 19 sera diluted with animal sera is added to the substrate. 20 The resultant mass is then incubated and washed. 21 Thereafter, radiolabeled, e.g., iodinated, e.g., I 125, 22 antibodies to human IgG or IgM is added to the mass. The 23 resultant mass is then washed and counted, e.g., in a 24 gamma-counter. If the count is higher than a count 25 performed on a normal sera control, the specimen contains 26 antibodies to the pre-S region of HBV. 27 It is believed that the above procedure for 28 detection of antibodies to the pre-S region of HBV can be 29

applied as a diagnostic tool in detection hepatitis B valus infection.

The pre-S protein moiety appears to be directly involved in attachment of HBV to liver cells of the host. Similar proteins are likely to be involved in the attachment of other viruses, the target of which is the liver. For this reason, synthetic peptides corresponding to the pre-S protein, as well as antibodies to them, could serve as the basis for diagnostic assays of and vaccines against other hepatitis viruses reacting with the same liver receptors as does hepatitis B virus.

In the above described procedures involving radioimmunoassay (RIA), an enzyme linked antibody can replace the radiolabeled antibody and ELISA techniques can be performed. Furthermore, fluorescence techniques can be employed in place of RIA or ELISA.

The labelling ("marking") of one of the reaction components can be brought about by use of a "marker" or "marker substance" such as by incorporation of a radioactive atom or group, or by coupling this component to an enzyme, a dyestuff, e.g., chromophoric moiety or a fluorescent group.

The components concerned are preferably labelled by coupling to an enzyme, since the estimation of this is much simpler than for example, the estimation of radioactivity, for which special apparatus is necessary.

The enzymes used are preferably those which can be colorimetrically, spectrophotometrically, or fluorimetrically determined. Non-limiting examples of enzymes for use in the present invention include enzymes

from the group of oxidoreductases, such as catalase, 2 peroxidase, glucose oxidase, beta-glucuronidase, beta-D-glucosidase, beta-D-galactosidase, urease and galactose oxidase. 5 The coupling of the enzyme and the immunological 6 component can be brought about in a known way, for example, 7 by the formation of an amide linkage by methods known from peptide chemistry. 9 The labelling with a radioactive isotope can also 10 be performed in a known way. Isotopes useful for labelling 11 are predominantly  $I^{125}$ ,  $I^{131}$ ,  $C^{14}$ , and  $H^3$ . 12 The incubation steps utilized in carrying out the 13 above procedures can be effected in a known manner, such as 14 by incubating at temperatures of between about 20°C and 15 about 50°C for between about 1 hour and about 48 hours. 16 Washings as described above are typically effected 17 using an aqueous solution such as one buffered at a pH of 18 6-8, preferably at a pH of about 7, employing an isotonic 19 saline solution. 20 The present invention also concerns diagnostic 21 test kits for conducting the above-described methods for 22 detecting antigens and antibodies. 23 A diagnostic test kit according to the present 24 invention for detecting antigens coded for the pre-S gene of 25 HBV in a test sample, would include the following: 26 a solid substrate coated with antibodies to a 27 peptide having an amino acid chain corresponding to at least 28 six consecutive amino acids within the pre-S gene coded 29

region of the envelope of HBV, the peptide free of an amino

1	acid sequence corresponding : the naturally occurring
2	proteins of HBV,
3	b. a protein-containing solution to saturate
4	protein binding sites on the solid subtrate, and
5	c. a given amount of radiolabeled antibody, such
6	antibody to either the peptide or HBsAg.
7	A diagnostic test kit according to the present
8	invention for detecting antibodies to the pre-S region of
9	hepatitis B virus in a test sample, would include the
10	following:
11	a. a solid substrate having adsorbed thereon a
12	peptide having an amino acid chain corresponding to at least
13	six consecutive amino acids within the pre-S gene coded
14	region of the envelope of HBV, the peptide free of an amino
15	acid sequence corresponding to the naturally occurring
16	proteins of HBV, the substrate being exposed to a
17	protein-containing solution to saturate protein binding
18	sites on the solid substrate, and
19	b. a given amount of radiolabeled antibodies to
20	human IgG or IgM.
21	Radiolabeled antibodies used in the
22	above-described test kits can be packaged in either solution
23	form, or in lyophilized forms suitable for reconstitution.
24	In the above test kits, enzyme or fluorescent
25	labelled antibodies can be substituted for the described
26	
27	radiolabeled antibodies.
28	The above described process and test kit for
29	detection of antibodies to the pre-S region of hepatitis B
30	virus can be utilized in many applications, such as

	# <b>*</b> ***
1	(1) detecting HBV infection in a patient by
2	`taking serum from the patient and applying the above
3	described test or using the above described test kit; and
4	(2) predicting recovery from HBV infection by
5	taking serum from an infected patient and applying the above
6	described antibody detection procedures.
7	The above described test procedure and test kit
8	for antibody detection can be used for making qualitative
9	comparisons between different HBV vaccines by taking serum
LO	from vaccinated patients and then utilize the
Ll	above-described test procedure or kit for antibody
12	detection. In general all known immunoassays using this
13	antigen as reagent can be performed using the synthetic
1.4	peptide of this invention. Generally all known immunoassays
15	using antibody containing serum or reagents can be performed
16	using antibody serum produced through the use of a synthetic
17	peptide of this invention. These immunoassays included all
18	those disclosed by Langone and Van Vunakis, Methods of
19	Enzymology, Academic Press, Volumes 70, 73 and 74. Those
20	assays disclosed in the disclosures of the following U.S.
21 22	Patents: 4,459,359; 4,343,896; 4,331,761; 4,292,403;
23	4,228,240; 4,157,280; 4,152,411; 4,169,012; 4,016,043;
24	3,839,153; 3,654,090 and Re 31,006 and volumes 70, 73 and 74
25	of Methods of Enzymology are incorporated herein by
26	reference.
27	A hepatitis B vaccine can be prepared by directly
28	using a conjugate of a lipid vesicle and a peptide
29	containing an amino acid chain corresponding to at least six
30	consecutive amino acids within the pro-5 gone coded region

of the surface antigen of hepocitis B virus in an appropriate buffer. The conjugate having peptide in the appropriate concentration can be used as a vaccine with or without an adjuvant, such as, e.g., aluminum hydroxide or others.

The active component of the vaccine can be employed with a physiologically acceptable diluent (medium), e.g., phosphate buffered saline. Generally speaking, the synthetic peptide concentration in a physiologically acceptable medium will be between approximately less than 1 miligram and more than 10 micrograms per dose.

The vaccine can be prepared and used in the same general manner as disclosed in U.S.P. 4,118,479, the entire contents of which are incorporated by reference herein.

intradermal or intramuscular injection. While the preferred route would depend upon the particular vaccine, it is believed that intramuscular injection will be generally suitable. Frequency of administration will vary depending upon the vaccine. Generally speaking, the vaccine will be administered in two doses about one month apart followed by a booster at six months to one year after primary immunization. The subsequent doses or the booster will depend on the level of antibody in the blood as a result of the initial immunization, and in certain instances may be unnecessary.

The hepatitis vaccine of the present invention is recommended for all persons at risk of developing hepatitis B infection and particularly those at especially high risk

0154902 such as patients and staff on hemodialysis unit, n ical 2 . personnel, persons of tropical populations and those 3 visiting the tropics. In the case of tropical populations, 4 particularly in Africa, Asia, the Mediterranean region and South America, where high incidence of hepatitis B 6 infections has been consistently observed, the vaccine 7 should be administered sufficiently early in life to prevent R acquisition of chronic carrier state infection which tend to 9 occur in these regions within the first five years of life. 10 In fact, the vaccine is expected to be useful for all 11 persons not already protected against hepatitis B infections 12 as a result of prior immunity. 13 In order to more fully illustrate the nature of 14 the invention and the manner of practicing the same, the 15 following non-limiting examples are presented: 16 **EXAMPLES** 17 Example 1 18

SDS-Polyacrylamide Gel Electrophoresis Of HBsAg.

About 20 and 200 ug, respectively, of HBsAg were separately electrophoresed for silver staining and transfer to nitrocellulose, respectively. Before electrophoresis, HBsAg was treated for 30 minutes at 37°C with 2-mercaptoethanol and NaDodSO $_4$  (10 mg/ml each in 8 M urea, 0.0625 M Tris, pH 7.2). Similar results were obtained with HBsAg alkylated with iodoacetate after reduction. HBsAg was purified and radiolabeled as described (A.R. Neurath, N.

Strick, C.Y. Huang, Intervirology, 10, 265 (1978)).

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SDS-Polyacrylamide gel cleatrophoresis ("SDS-PAGE") was carried out following published procedures.

See V.K. Laemmli, Nature (London), 227, 680 (1970). 1 However, in order to maintain proteins in fully denaturated form, 8M urea was utilized in the running buffers in 3 electrophoresis. Polypaptides separated by SDS-PAGE were 5 transferred to nitrocellulose using the TE 42 Transphor unit 6 9 (Hoefer Scientific Instruments, San Francisco, California) 7 following the procedure recommended by the manufacturer. The 8 transferred proteins were tested for determinants reacting 9 with antibodies to intact HBsAg (anti-HBs) using 10  $^{125}$ I-labeled human anti-HBs supplied as part of a commercial 11 test kit (Abbott Laboratories, North Chicago, Illinois) as 12 13 described (J.C. McMichael, L.M. Greisiger, L. Millman, J. 14 Immunol. Meth., 45, 79, (1981)). 15 From the 20ug sample gel, separated HBsAg 16 polypeptides (their  $M_{_{\mbox{\scriptsize T}}}$  given in kilodaltons) were stained by 17 silver in situ (J.H. Morrissey, Anal. Biochem, 117, 307, 18 (1981)), (see Fig. 1, Panel a) to yield two major and 19 several minor polypeptides as expected. The separated 20 polypeptides from the other 200  $\mu\mathrm{g}$  sample gel was then 21 electrophoretically transferred to nitrocellulose, reacted 22 (probed) with 125I-labeled antibodies to intact HBsAg (anti 23 HBs) and submitted to autoradiography (Fig. 1b). 24 Surprisingly, the 33 and 36 kilodalton (P33 and 25 P36), rather than the two most abundant polypeptides reacted 26 preferentially with anti-HBs (Fig. 1, Panel b). This 27 suggested the presence of disulfide bond independent 28 antigenic determinants reacting with anti-HBs on amino acid 29 sequences which are not coded for by the S-gene of HBV DNA.

P33 and P36 ontain the sequence corresponding to the product of the S-gene and additional 55 residues at the amino-terminal part starting with Met at position 120 in the pre-S gene region (See Fig. 2).

#### Example 2

# Synthesis Of A Peptide Mimicking Antigenic Determinants Corresponding To Residues 120-145 Of The Pre-S Gene Product

The location of antigenic determinants on proteins may be predicted from computing the relative hydrophilicity along the amino acid sequence. See T.P. Hopp, K.R. Woods, Proc. Natl. Acad. Sci. USA, 78, 3824 (1981) and J. Kyte, R.F. Doolittle, J. Mol. Biol., 157, 105 (1982).

Results of such computation (J. Kyte et al supra) for the translation product of the pre-S region are shown in Fig. 3 and suggest the location of antigenic determinants in the sequence to the right from Met 120 within residues 120-140. The segment corresponding to residues 120-145 (Fig. 2) (pre-S 120-145, subtype adw<sub>2</sub>) was selected for synthesis.

A C-terminal Cys(-SH containing) residue was added to allow unambiguous conjugation to carrier molecules and affinity matrices, while leaving the N-terminal unblocked as it may be in the intact protein. The molecule contains one Tyr and can therefore be radiolabeled. The Tyr could also be used for conjugation, although it might be a part of the antigenic determinant.

The peptide was synthesized by an accelerated version of stepwise solid phase peptide synthesis on the benzhydrylamine-type resin of Gaehde and Matsueda (Int. J.

1	Peptide Protei. Res., 18, 451, (1981)) using
2	Boc-NH-CH(phenyl)-phenyl-OCH2COOH to derivatize NH2CH2-Resin
3	'(A.R. Mitchell, S.B.H. Kent, M. Engelhard and R.B.
4	Merrifield, J. Org. Chem., 43, 2845-2852, (1978)). After the
5	Cys was coupled, the protected peptide chain was assembled
6	according to the following protocol:
7	1. Deprotection: 65% v/v trifluoroacetic acid in
8	dichloromethane, 1x10 minutes;
9	2. Wash: a flowing stream of dichloromethane was
10	run over the resin under suction from an aspirator for 20
11	seconds;
12	3. Neutralization: 10% v/v diisopropylethylamine
13	in dichloromethane, 2x1 minutes;
14	4. Wash: a flowing stream of dichloromethane was
15	run over the resin under suction from as aspirator for 20
16	seconds;
17	5. Coupling: 2 mmol tert.Boc-L-amino acid in 2ml
18	dichloromethane was added to the neutralized resin followed
19	immediately by 1mmol dicyclohexylcarbodiimide in 2ml
20	dichloromethane; after 10 minutes a sample of resin
21	(approximately 5mg) was taken for determination of coupling
22	yield by quantitative ninhydrin, and 10ml dimethylformamide
23	was added and the coupling continued. (Asn and Gln were
24	coupled in the presence of hydroxybenzotriazole).
25	6. After the ninhydrin determination of a
26	satisfactory coupling, the resin was washed as in step 4,
27	above. For the addition of subsequent residues, the cycle
28	
29	was repeated. If recoupling was necessary, steps 3-5 were

repeated. The synthesis was performed on a 0.5mmol scale

were 10ml except where noted.

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Protected amino acid derivatives used were N-alpha-tert.butyloxycarbonyl protected and side chain protected as follows: Arg(NGTosyl); Cys(4MeBzl); Tyr(BrZ); Asp(OBzl); Thr(Bzl); His(ImTosyl). Met and Trp were unprotected on the side chains. In another synthesis, otherwise identical, use of His(ImDNP) and Trp(InFormyl) gave purer product.

Assembly of the peptide chain was monitored by the quantitative ninhydrin reaction (V.K. Sarin, S.B.H. Kent, J.P.Tam, R.B. Merrifield, Anal. Biochem, 117, 147-157, (1981)) and was without difficulty except for the addition of the histidine residue which was 10% incomplete despite repeated couplings, presumably due to an impure amino acid derivative. After assembly of the protected peptide chain, the N-terminal Boc group was removed by trifluoroacetic acid treatment and the resin neutralized as in steps 1-4 above. Then the peptide was cleaved and deprotected by a 1 hour treatment at 0°C with HF containing 5% v/v p-cresol and 5% v/v p-thiocresol to give the desired peptide as the C-terminal cysteinamide. Where His(ImDNP) was used, the DNP was removed by treatment with phenylphenol prior to HF cleavage. Where TrP (InFormyl) was used, HF conditions were adjusted to remove the Formyl group; either HF containing 10% anisole and 5% 1,4-butanedithiol, or HF containing p-cresol and 5% 1,4-butanedithiol. The product was precipitated and washed by the addition of ether, then dissolved in 5% v/v acetic acid in water and lyophilized to

1	give a fluffy white soll(
2	Quantitative Edman degradation (H.D. Niall, G.W.
3	Tregear, J. Jacobs, Chemistry and Biology of Peptides, J.
4	Meienhofer, Ed (Ann Arbor Press, Ann Arbor, MI, 1972), pp.
5	659-699) of the assembled peptide-resin revealed a high
6	efficiency of chain assembly (S.B.H. Kent, M. Riemen, M.
7	LeDoux, R.B. Merrifield, Proceedings of the Fourth
8	International Symposium on Methods in Protein Sequence
9	Analysis, M. Elzinga, Ed. (Humana, Clifton, New Jersey,
10	1982), pp. 626-628) which proceeded at a $\geq$ 9./9.7 percent
11	efficiency at each step, except for histidine at sequence
12	position pre-S 128. HPLC of the peptide cleaved off the
13	resin revealed a single major peak corresponding to
14	approximately 85 percent of peptide material absorbing light
15	at 225 nm.
16	Examples 3-6
17	Immunologic Properties Of A Peptide Mimicking
18	Antigenic Determinants Corresponding To Residues 120-145 of
19	the Pre-S Gene Product (pre-S 120-145)
20	CHC 123
21	Example 3
22	Immunization
23	Immunization of rabbits with either free or
24	carrier-bound pre-S 120-145 (subtype adw <sub>2</sub> ) were conducted
25	carrier-bound pre-5 120-145 (Substituting 2)
26	and resulted in an antibody response in all animals against
27	both the homologous peptide and HBsAg (Fig. 4).
28	The peptide corresponding to the amino acid
29	sequence 120-145 (pre-S 120-145) of the pre-S region of HBV
20	DNA (subtype adwa; P. Valenzuela, P. Gray, M. Quiroga, J.

Zaldivar H.M. Goodman, W.J. Rutter, Nature (London), 280, 815, (1979)) containing an additional Cys residue at the C-terminal, added for convenience of coupling to carriers, was synthesized by an improved solid phase technique (S.B.H. Kent, Biomedical Polymers, E.P. Goldberg, A. Nakajima, Eds. (Academic, New York, 1980), pp. 213-242; A.R. Mitchell, S.B.H. Kent, M. Engelhard, R.B. Merrifield, J. Org. Chem. 43, 2845, (1978); and S. Mojsov, A.R. Mitchell, R.B. Merrifield, J. Org. Chem., 45, 555 (1980).

For immunoassays and linking to carriers the peptide was treated with 2-mercaptoethanol and separated

For immunoassays and linking to carriers the peptide was treated with 2-mercaptoethanol and separated from low Mr components by chromatography on Sephadex G-10 (A.R. Neurath, S.B.H. Kent, N. Strick, Proc. Natl. Acad. Sci. USA, 79, 7871 (1982)).

Groups of two to three rabbits were immunized with either free pre-S 120-145 or with the peptide linked to cysteine-activated liposomes containing stearylamine, dilauroyl lecithin and cholesterol which had been fixed with glutaraldehyde, and either did or did not have attached RAT groups for enhancing antibody responses to haptens (A.R. Neurath, S.B.H. Kent, N. Strick, J. Gen. Virol., in press (1984)). The immunization schedule involving the use of complete and incomplete Freund's adjuvant was the same as described (Neurath, Kent, Strick, et al (1984) supra). Antibodies to HBsAg in sera of rabbits immunized with pre-S 120-145 were tested by a double-antibody radioimmunoassay (RIA) using HBsAg-coated polystyrene beads and 125I-labeled anti-rabbit IgG (Neurath, Kent, Strick, et al (1984) supra).

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l	Antibodies to the homologous peptide were tested \
2	by a similar test except that 2.5 mg of a cellulose-peptide
3	conjugate were used instead of coated beads. This conjugate
4	was prepare in the following way: 0.5 g of sulfhydryl
5	cellulose, prepared as described (P.L. Feist, K.J. Danna,
6	Biochemistry, 20, 4243 (1981)), were suspended in 5 ml 0.1 M
7	sodium acetate, pH 5, and mixed with 2.5 ml of 0.25 M
8	N-N'-p-phenylenedimaleimide in dimethylformamide for one
9	hour at 30°C and then washed with 0.1 M phosphate-10mM EDTA,
10	pH 7.0. The cellulose derivative was suspended in 10 ml of
11	the latter buffer containing 5 mg of pre-S 120-145 and mixed
12	for at least sixteen hours at 20°C. The cellulose derivative
13	was extensively washed and suspended in 0.14 M NaCl-10 mM
14	Tris-3 mM NaN3 (TS). The final preparation contained 8 mg of
15	pre-S 120-145 per g of cellulose.
16	
17	Example 4
18	Radioimmunioassays were conducted with several
19	dilutions of a serum from one of the rabbits immunized with
20	pre-S 120-145 linked to liposomes (See Fig. 5).
21	Antibodies were still detectable when the antisera
22	were diluted up to 1.6 $\times$ 10 <sup>5</sup> -fold (Fig. 5).
23	Pre-S 120-145 or anti-pre-S 120-145 inhibited the
24	reaction between $^{125}$ I-labeled anti-HBs and P33 (P36).
25	125 <sub>I-labeled HBsAg was immunoprecipitated with anti-pre-S</sub>
26	120-145 at all dilutions positive by RIA (Fig. 5). HBV
27	particles reacted with anti-pre-S 120-145 as determined by
28	detection of HBV-DNA within the immune complexes and by
29	detection of uparami and

1	electron microscopy (A.R. Neurath, N. Strick, L. Baker, S.
2	Krugman, Proc. Nat. Acad. Sci. USA, 79, 4415 (1982)).
3	
4	Example 5
5	Anti-Peptide Antibody as A Specific Probe for
6	Detection of P33 and P36
7	Anti-pre-S 120-145 was reacted with P33 and P36.
8	HBsAg polypeptides separated by SDS-PAGE run in urea were
9	
10	transferred to nitrocellulose, reacted with anti-pre-S
11	120-145 diluted 1/80 in TS containing 10 mg/ml of bovine
12	serum albumin and 2.5 mg/ml of gelatine (TS-BG) for five
13	hours at 20°C. To detect bound IgG, the nitrocellulose sheet
14	was washed and exposed to $^{125}$ I-labeled protein A (0.4 $\mu$ C/100
15	ml TS-BG) for five hours at 20°C. For further details see
16	Fig. 1. In Fig. 6, arrows indicate the positions of P33 and
17	P36. The top arrow (corresponding to a molecular weight of
18	66 kilodaltons) indicates another protein reacting with
19	anti-pre-S 120-145, possibly corresponding to a dimer of
20	P33.
21	
22	Example 6
23	Development Of A Diagnostic Test For The
24	Detection Of Antigens Coded For By The Pre-S Gene In Sera
25	Of HBV-Infected Individuals
26	Fig. 7 shows the results of a diagnostic test
27	based on polystyrene beads coated with anti-pre-S 120-145.
28	Serial dilutions of an HBsAg-positive serum in a
29	mixture of normal human and rabbit serum each diluted 1/10
30	in TS were tested. 125 I-labeled human anti-HBs (Abbott

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ı	Laboratories)as used in the test performed essentially as
2	described for the AUSRIA II diagnostic kit (Abbott
3	Laboratories). Results are expressed as RIA ratio units,
4	determined by dividing cpm corresponding to positive samples
5	by cpm corresponding to positive samples by cpm
6	corresponding to normal serum controls. The endpoint titer
7	corresponding to normal trace corresponds to the highest dilution at which the RIA ratio
8	was 2.1 (broken line). The endpoint titer of the serum as
	was 2.1 (broken line). The start was approximately 1/10 <sup>6</sup> .
9	determined by the AUSRIA test was approximately 1/10 <sup>6</sup> .
10	Negative results were obtained with control beads coated
11	with normal rabbit IgG.
12	Similar results were obtained with sera containing
13	HBsAg subtypes ad and ay, indicating that the synthetic
14	peptide with the sequence corresponding to subtype adw (Fig.
15	2) carried common group-specific antigenic determinants.
16	2, 000
17	Example 7
18	Synthesizing and Testing
19	S(135-155) Derivatives
20	Each of the conjugates ((1) to (26)) of S(135-155)
21	listed in Table 1, except conjugate 3, was mixed 1:1 with
22	complete Freund's adjuvant and injected into two New Zealanc
23	White rabbits (65 to 160 µg of peptide per rabbit). The
24	rabbits were further injected at biweekly intervals with
25	equal doses of conjugates in incomplete Freund's adjuvant
26	(not used for conjugate 3). Blood specimens were taken two
27	weeks after each injection.
28	To prepare conjugates 1, and 4-8 (Table 1), 1 mg
29	quantities of peptide 309-329 of the env gene product
	(S(135-155)) were activated with a two times molar excess o

1 N-ethyl-N'(dimethyl-aminopropyl) carbodiimide (EDAC) and 2 N-hydroxy-benzotriazole (NHBTA) and subsequently linked to 3 equimolar quantities of poly-D-lysine and diaminoalkanes 4 (from Fluka AG, Buchs, Switzerland), respectively, as 5 described (Arnon, R., Sela, M., Parant, M. and Chedid., L., "Antiviral Response Elicited By A Completely Synthetic 7 Antigen With Built-In Adjuvanticity", Proceedings of The National Academy of Science USA, 77,6769-6772, (1980)). To 9 prepare conjugates 2 and 3, 1 mg quantities of each 10 EDAC-activated S(135-155) and MDP (Calbiochem, San Diego, 11 California) were linked to 10 mg of poly-D-lysine. Peptide 12 309-329 of the env gene product (800µg) was oxidized with 13 ferricyanide (Dreesman et al, 1982 supra), activated with 14 EDAC as above and linked to 4 mg of LPH. Chromatography on 15 Sephadex G-25 indicated complete linking of the peptide to 16 LPH (conjugate 9). The oxidized, EDAC-activated peptide (1 17 mg) was also conjugated to 1 mg of polyvaline in a suspension of 2.5 ml of 1 M NaHCO3, pH 8.5, and 10 ml of 19 CHCl3. The interphase and aqueous phase after centrifugation 20 was used for immunization (conjugate 10). 21 Liposomes were prepared by the method of Oku, N. 22 Scheerer, J.F., and MacDonald, R.C., "Preparation of Giant 23

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Liposomes", Biochimica et Biophysica Acta, 692, 384-388 (1982). Stearylamine, dilauroyl lecithin and cholesterol were dissolved in glucose-saturated ethanol at final concentrations of 10, 23 and 1.43 mg/ml, respectively. For some liposome preparations, the concentration of dilauroyllecithin was decreased to 17.5 mg/ml and sphingomyelin was added (10 mg/ml). Other preparations contained as an

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additional component lipid A (420µg/ml; Calbiochem). The solutions were dialyzed against 0.1 M NaCHO3, pH 8.5, in dialysis bags with a molecular weight cut-off of 103 daltons for at least sixteen hours. The liposomes were treated for approximately six hours with glutaraldehyde (final concentration 30 mg/ml), mixed with 0.5 volumes of 33.9% (w/w) sodium diatrizoate, floated four times into 1 M NaCHO, by centrifugation for ten minutes at 10,000 rpm, and reacted with 0.84 to 1 mg of peptide 309-329 of the env gene product per 10 mg stearylamine overnight at 20°C. The linking of peptide 309-329 of the env gene product to liposomes under these conditions was complete. Some preparations were reacted additionally with 7.5 mg of RAT (Biosearch, San Rafael, California) per 10 mg of stearylamine for six hours at 20°C. The liposomes were floated three times into 0.14 M NaCl, 0.01 Tris-HCl-0.02% NaN, (TS) and dialyzed against TS-10<sup>-4</sup> M oxidized glutathione for at least sixteen hours.

In some cases (20) and (21) the stearylamine-containing liposomes were not derivatized with GA but instead directly reacted with EDAC-activated peptide 309-329 of env gene product. Alternately, (18) and (19), the activated peptide 309-329 of env gene product was linked to glutaraldehyde-treated liposomes further derivatized by reaction with 0.2 M ethylene diamine at pH 8.5 overnight at 20°C followed by floating two times into 0.1 M NaHCO<sub>3</sub>, pH 8.5, reduction with 10uM sodium dithionite for one hour at 20°C and repeated floating into the same buffer. An aliquot of these liposomes was additionally reacted with

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EDAC-activated RAT. The liposomes were finally dialyzed against TS-10<sup>-4</sup> M oxidized glutathione.

In one preparation (22), stearic acid was used instead of stearylamine for the preparation of liposomes. These were dialyzed against 0.01 M NaCl, activated with EDAC (50 mg/ml for two hours plus additional 25 mg/ml for one hour) at pH 5.5 and 20°C, floated two times into 0.01 M NaCl and reacted with the peptide 308-329 of the env gene product in 1 M NaHCO<sub>3</sub>, pH 8.5, overnight.

Polyglutaraldehyde microspheres were prepared as described by Margel, S., Zisblatt, S. and Rembaum, A. "Polyglutaraldehyde: A New Reagent For Coupling Proteins To Microspheres And For Labeling Cell-Surface Receptors. II. Simplified Labeling Method By Means Of Non-Magnetic And Magnetic Polyglutaraldehyde Microspheres\*, Journal of Immunological Methods, 28, 341-353 (1979), using Polysurf 10-36 B (Bartig Industries Inc., New Canaan, Conn., Margel & Offarim, (1983)). One mg of the peptide 309-329 of the env gene product was linked to approximately 50 mg of microspheres under conditions similar to those described for glutaralde-hyde treated liposomes. Conjugate 25 was prepared ~ by treating the microspheres with 5 ml of 0.1 ME-amino caproic acid at pH 8.5 overnight. After centrifugation, the microspheres were suspended in dimethylformamide (2ml) and reacted with 2 mg EDAC plus 670 ug NHBTA for one hour at 20°C. After centrifugation, the microspheres were resuspended in 2 ml of 0.1 M NaHCO3, pH 8.5, containing 1 mg of peptide 309-329 of the env gene product.

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All reagents listed above were of analytical grade 2 and obtained from Sigma, St. Louis, Missouri, unless 3 indicated otherwise. Free peptide 309-329 of the env gene product (mol. 5 weight = 2,664 daltons) containing five cysteine residues 6 was in a predominantly monomeric form, since it was eluted after molecular exclusion chromatography in about the same 8 fractions as insulin A chain. Linking to diaminobutane and 9 to other diamino-alkanes (data not shown) resulted in 10 formation of S(135-155) polymers which were immunogenic and 11 induced both antipeptide and anti-HBs antibodies. 12 Preparations (4), (5) and (7) also induced anti-HBs, while 13 polymers with diaminooctane or dodecane linkers (6) and (8) 14 failed to do so (Fig. 8) for reasons not known. Oxidation of 15 the peptide 309-329 of the env gene product resulted in 16 polymerization (data not shown). The polymer linked to LPH 17 (conjugate 9) induced high levels of anti-S(135-155) but no 18 anti-HBs, unlike S(135-155) linked to KLH or LPH in its 19 reduced form (Neurath et al., 1982, supra). This finding 20 again emphasizes the role of peptide conformation in 21 inducing antibodies to the native protein. Linking of the 22 oxidized peptide to highly hydrophobic poly-L-valine 23 resulted in a conjugate (10) of low immunogenicity. 24 S(135-155) linked to poly-D-lysine administered with 25 Freund's adjuvant (1) or having covalently linked MDP and 26 given without adjuvant (3) induced both anti-S(135-155) and 27 anti-HBs. The latter conjugate administered with Freund's 28 adjuvant (2) appeared poorly immunogenic. S(135-155) linked 29 to glutaraldehyde treated liposomes containing stearylamine 30

1 (conjugate 11) induced levels of anti-HBs comparable to 2 those elicited by those elicited by conjugates with KLH or 3 LPH (Neurath et al., 1982, supra). Incorporation of sphingomyelin and.or lipid A, components reported to enhance 5 the antigenicity of haptens inserted into liposomal 6 membranes (Yasuda, T., Dancey, G.F. and Kinsky, S.C., 7 "Immunogenicity Of Liposomal Model Membranes In Mice: 8 Dependence On Phospholipid Composition", Proceedings Of The 9 National Academy Of Sciences, 74, 1234-1236 (1977)), into 10 the liposomes (conjugates 13, 15a, 16) failed to enhance 11 anti-HBs, responses. 12 Conjugates (18 and 19) prepared by linking 13 S(135-155) to glutaraldehyde-treated liposomes through an 14 ethylenediamine bridge rather than directly, had the 15 capacity to induce anti-HBs but a considerable variability 16 in response between individual rabbits was observed. 17 S(135-155) before or after oxidation and 18 subsequently linked to stearyl-amine-containing liposomes 19 (not fixed with glutaraldehyde; preparations 20 and 21) or 20 to stearic acid-containing liposomes (22) induced low levels 21 of anti-S-135-155 and no measurable anti-HBs. 22 S(135-155) linked directly to microspheres of 23 polyglutaraldehyde (preparations 23 and 24) induced a 24 primary anti-HBs response. However, the level of anti-HBs 25 decreased in the course of immunization. Anti-HBs was un-26 detectable in sera collected two weeks after the third 27 immunization. S(135-155) linked to these microspheres 28 through & -amino-caproic acid (25) and 1-cysteine (26) 29

bridges, respectively, either failed (25) or was marginally efficient (26) in eliciting anti-HBs.

s(135-155)-KLH or LPH conjugates elicited a primary anti-HBs response but the level of anti-HBs failed to increase in sera of rabbits after additional antigen doses (Neurath et al., 1982 <a href="mailto:supra">supra</a>). With the conjugates described above, generally, a decrease of anti-HBs levels was observed four or six weeks after primary immunization (Fig. 9B), but exceptions were observed in a minority of rabbits (panel 5, Fig. 9A). This declining trend was uniformly reversed when RAT was inserted into liposomal membranes together with S(135-155) (for example Fig. 9C and Fig. 9D).

The immunogenicity of haptens inserted into liposomal membranes depends on the phospholipid composition of the liposomes and seemed to be inversely related to the fluidity of these membranes (Yasuda et al., 1977 <a href="supra">supra</a>; Dancey, G.F., Yasuda, T. and Kinsky, S.C., "Effect Of Liposomal Model Membrane Composition On Immunogenicity", The Journal Of Immunology, 120, 1109-1113 (1978)).

Treatment of stearylamine-containing liposomes with glutaraldehyde was found to provide reactive groups suitable for linking of synthetic peptides and at the same time increases the rigidity of the lipid membranes. Such liposomes, especially when containing carrier function enhancing RAT sites (Alkan, S.S., Nitecki, D.E. and Goodman, J.W., "Antigen Recognition And the Immune Response: The Capacity of 1-Tryosine-Azobenzenearsonate To Serve As A Carrier For A Macromolecular Hapten", The Journal Of

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        Immunology, 107, 353-358, (1971), and Alkan, S.S., Williams,
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        E.B., Nitecki D.E. and Goodman, J.W.. "Antigen Recognition
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        And the Immune Response. Hurmoral And Cellular Immune
4
        Responses To Small Mono- And Bifunctional Antigen
5
        Molecules", The Journal Of Experimental Medicine, 135,
6
        1228-1246, (1972)), are a promising tool for preparing fully
7
        synthetic immunogens for eliciting anti-viral antibodies.
8
9
                                     TABLE 1
10
                   List of cross-linkers and carriers used
                 for the preparation of S(135-155) conjugates
11
12
                   Poly-D-lysine (mol. weight 3-7 x 10^4)
13
         (1)
         (2)
                   1 + N-Acetylmuramyl-L-alanyl-D-isoglutamine (MDP)
14
15
         (3)
                   = 2
16
                   1,4-diaminobutane
         (4)
17
                   1,6-diaminohexane
         (5)
18
         (6)
                   1,8-diaminooctane
19
         (7)
                   1,10-diaminodecane
20
         (8)
                   1,12-diaminododecane
21
                   Oxidized S(135-155) linked to LPH
         (9)
22
                   Oxidized S(135-155) linked to poly-L-valine
         (10)
23
                   Liposomes containing stearylamine, and treated
         (11)
                   with glutaraldehyde
24
                   = 11 = L-tyrosine-azobenzene-p-arsonate (RAT)
         (12)
25
         (13)
                   = 11 + Sphingomyelin (from bovine brain)
26
                   = 13 + RAT
         (14)
27
                   = 11 + Lipid A
         (15a)
28
         (15)
                   = 15a + RAT
29
30
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1
         (16)
                   = 13
                         + Lipid A
 2
         (17)
                   = 16
                         + RAT
 3
         (18)
                   = 11 treated with ethylenediamine
         (19)
                   = 18 + RAT
 5
         (20)
                   = Liposomes containing stearylamine reacted
                     with oxidized S(135-155) (see 9)
 6
         (21)
                   = 20 except S(135-155) was oxidized after
 7
                     attachment to liposomes
 8
         (22)
                   Stearic acid containing liposomes
 9
         (23)
                   Polyglutaraldehyde micropheres
10
         (24)
                   = 23 + RAT
11
         (25)
                   = 23 treated with \mathcal{E}-aminocaproic acid
12
         (26)
                   = 23 treated with L-cysteine
13
14
        Example 8
15
                   A peptide pre-S (12-32) (subtype adw<sub>2</sub>) was
16
        synthesized according to the procedure described hereinabove
17
         in Example 2.
                        The free peptide, the peptide linked to
18
        glutaraldehyde cross-linked liposomes (±RAT groups)
19
         (according to the procedure described above in Example 7) as
20
        well as the peptide linked to KLH were used to immunize
21
        rabbits. The corresponding antibodies recognized not only
22
        the peptide, but also HBsAg and HBV. In view of the above,
23
        this peptide is believed quite useful for a vaccine against
24
        hepatitis B virus, and as the basis of useful HBV
25
        diagnostics based on either the peptide itself (to detect
26
        anti-HBV response in infected or immunized individuals), or
27
        on peptide antibodies to detect hepatitis B antigens.
28
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1 Example 9

A peptide pre-S (117-134) (subtype adw<sub>2</sub>) was

synthesized according to the procedure described hereinabove in Example 2.

#### Example 10

A rabbit was immunized with the peptide pre-S (117-134) prepared according to Example 9 and linked to a carrier according to the procedure of Example 7. Such immunization was conducted according to the procedure described hereinabove in Example 3 and was found to produce antibodies in the serum of the rabbit so innoculated. However, the antibody titers were substantially less than those observed for the use of pre-S (120-145) and pre-S (12-32).

#### Example 11

The immune response in rabbits to each of two synthetic peptides corresponding to residues 120-145 and 12-32 of the translational product of the pre-S gene of HBV DNA (subtype adw<sub>2</sub>) was tested. Peptide pre-S (120-145) was prepared according to Example 2 and peptide pre-S (12-32) was prepared according to Example 8. Their sequences are: MOWNSTAFHQTLQDPRVRGLYLPAGG (pre-S (120-145)) and MGTNLSVPNPLGFFPDHQLDP (pre-S (12-32)). For immunization, the peptides were used in free form, employing alum or Freund's adjuvant, or linked to carriers, i.e., keyhole lympet hemocyanin (KLH) and cross-linked liposomes, respectively. The liposomes were prepared as described in

Example 7.

The best results were obtained with peptides covalently linked to the surface of liposomes (see Fig.10). Immunization with KLH conjugates resulted in a high anti-KLH response (endpoint titers of 1/5,000,000 by radio-immunoassay), apparently causing low booster responses to the peptides. On the other hand, much lower antibody responses (approximately 1/10<sup>3</sup>) to RAT groups were detected, when RAT-containing liposomes were used as carriers. Antibodies to liposomes (lacking RAT) were undetectable. This suggests that liposomes are the carrier of choice for immunization with synthetic peptides.

#### Example 12

To establish whether or not antigenic determinants corresponding to pre-S gene coded sequences are preferentially present on HBV particles, the reaction of antisera raised against HBV particles with the two synthetic peptides analogues of the pre-S protein was tested. The maxium dilutions of this antiserum at which antibodies reacting with the synthetic peptides were still detectable were: approximately 1/62,500 ( $1/2 \times 10^6$  with tests utilizing 125 I-labeled protein A instead of labeled second antibodies), and approximately 1/2,560 for peptides pre-S(120-145) and pre-S(12-32), respectively (see Fig. 11). The antiserum (adsorbed on HBsAq-Sepharose to remove antibodies to S-protein) did not react with synthetic peptide analogues of the S-protain, peptide (309-329) of the env gene product (S(135-155)), peptide (222-239) of the env gene product (S(48-65)) and peptide (243-253) of the env

gene product (S(69-79)) and was, therefore, specific for 2 pre-S gene coded sequences. In comparison, the dilution 3 endpoints of antisera prepared against the homologous peptides were approximately 1/300,000 and approximately 5 1/80,000 for anti-pre-S(120-145) (see Fig. 11) and 6 anti-pre-S(12-32) (data not shown), respectively. 7 The synthetic peptides were recognized also by 8 antibodies (IgG and IgM) in sera of individuals who had just 9 recovered from acute hepatitis B, and by rabbit antibodies 10 against a fusion protein between chloramphenicol 11 acetyltransferase and a portion of pre-S protein expressed 12 in E. coli (see Fig. 11). 13 On the other hand, humans vaccinated with 14 pepsin-treaded HBsAg (M.R. Hilleman, E.B. Buynak, W.J. 15 McAleer, A.A. McLean, P.J. Provost, A.A. Tytell, in Viral 16 Hepatitis, 1981 International Symposium, W. Szmuness, H.J. 17 Alter, J.E. Maynard, Eds. (Franklin Institute Press, 18 Philadelphia, PA, 1982), pp. 385-397) or with HBsAg produce 19 in yeast (devoid of pre-S gene coded sequences; W.J. 20 McAleer, E.B. Buynak, R.F. Maigetter, D.E. Wambler, W.J. 21 Milbur, M.R. Hilleman, Nature (London), 307, 178 (1984)) die 22 not develop detectable antibodies recognizing either of the 23 two synthetic peptides. On the other hand, 7 out of 12 24 individuals who received a vaccine consisting of intact 25 HBsAg developed these antibodies. 26 27 Example 13 28 Quantitative aspects of the immunological cross-29

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reactivity between pre-S gene coded sequences exposed on HB'

1 particles (or on HBsAg) and the synthetic peptide analogues 2 were tested. The peptides were conjugated to 3  $\beta$ -galactosidase, and the inhibitory effect of free peptides, HBV and HBsAg, respectively, on the formation of immune complexes containing the enzyme-conjugated peptide was studied. Results shown in Fig. 12 indicate that HBV. at sufficient concentrations, inhibited completely the reaction between anti-pre-S(120-145) and pre-S(120-145)-9  $\beta$ -galactosidase. HBsAg had < 1/5 of the inhibitory activity 10 corresponding to HBV. The inhibitory activity of 11 pepsin-treated HBsAg was < 1/1,000 of the activity 12 corresponding to intact HBsAg. These results indicate the 13 absence in the anti-pre-S(120-145) serum of a subpopulation 14 of antibodies which recognize the synthetic peptide but not 15 the native protein. Such antibody subpopulations are 16 observed in many other antisera raised against synthetic 17 peptide analogues of viral proteins. The concentration of 18 free peptide sufficient for approximately 50% inhibition of 19 the reaction of pre-S(120-145)- $\beta$ -galactosidase with 20 anti-pre-S(120-145) is approximately 1/100 of that for HBV 21 on a weight basis (see Fig. 11). However, since the 22 molecular weight of pre-S(120-145) is approximately 3 kD and 23 the molecular weight of HBV protein components reacting with 24 anti-pre-S(120-145) (representing a minor (< 20%) portion of 25 the total HBV mass) is between approximately 33 and 26 approximately 67 kD, the molar concentrations of HBV and 27 pre-S(120-145) required for this degree of inhibition are 28 approximately the same. This indicates that the antigenic 29 determinants on the peptide analogue and on the 30

1	corresponding segment of the HBV envelope protein(s) are
2	structurally closely related.
3 .	
4	Example 14
5	A peptide pre-S (94-117) (subtype adw <sub>2</sub> ) was
6	synthesized according to the procedure described hereinabove
7	in Example 2.
8	
9	Example 15
10	A rabbit was immunized with the peptide pre-S
11	(94-117) prepared according to Example 14 and linked to a
12	carrier according to the procedure of Example 7. Such
13	immunization was conducted according to the procedure
14	described hereinabove for Example 3 and was found to produce
15	antibodies in the serum of the rabbit so inoculated.
16	
17	However, the antibody titers were substantially less than
18	those observed for the use of pre-S (120-145) and pre-S
19	(12-32).
20	
21	Example 16
22	A peptide pre-S (153-171) (subtype adw <sub>2</sub> ) was
23	synthesized according to the procedure described hereinabove
24	in Example 2.
25	
26	Example 17
27	A rabbit was immunized with the peptide pre-S
28	(153-171) prepared according to Example 16 and linked to a
29	carrier according to the procedure of Example 7. Such
30	immunization was conducted according to the procedure

described hereinabove for Example 3 and was found to produce 2 antibodies in the serum of the rabbit so innoculated. 3 However, the antibody titers were substantially less than those observed for the use of pre-S (120-145) and pre-S 5 (12-32). 6 7 Example 18 8 A peptide pre-S (1-21) (subtype adw<sub>2</sub>) was 9 synthesized according to the procedure described hereinabove 10 in Example 2. 11 12 Example 19 13 A rabbit was immunized with the peptide pre-S 14 (1-21) prepared according to Example 18 and linked to a 15 carrier according to the procedure of Example 7. Such 16 immunization was conducted according to the procedure 17 described hereinabove for Example 3 and was found to produce 18 antibodies in the serum of the rabbit so innoculated. 19 However, the antibody titers were substantially less than 20 those observed for the use of pre-S (120-145) and pre-S 21 (12-32).22 23 Example 20 24 A peptide pre-S (32-53) (subtype adw<sub>2</sub>) was 25 synthesized according to the procedure described hereinabove in Example 2. 27 28

Example 21

A rabbit was immunized with the peptide pre-S (32-53) prepared according to Example 20 and linked to a carrier according to the procedure of Example 7. Such immunization was conducted according to the procedure described hereinabove for Example 3 and was found to produce antibodies in the serum of the rabbit so innoculated. However, the antibody titers were substantially less than those observed for the use of pre-S (120-145) and pre-S (12-32).

#### Example 22

A peptide pre-S (57-73) (subtype  $adw_2$ ) was synthesized according to the procedure described hereinabove in Example 2.

#### Example 23

A rabbit was immunized with the peptide pre-S (57-73) prepared according to Example 22 and linked to a carrier according to the procedure of Example 7. Such immunization was conducted according to the procedure described hereinabove for Example 3 and was found to produce antibodies in the serum of the rabbit so innoculated. However, the antibody titers were substantially less than those observed for the use of pre-S (120-145) and pre-S (12-32).

1 Example 24 2 Detection of anti-pre-S protein antibodies in 3 human sera using synthetic peptides. As discussed above, antibodies recognizing 5 synthetic peptide analogues of the pre-S protein were 6 detected in sera of humans during recovery from hepatitis B 7 (Fig. 11). The time course of development of antibodies 8 recognizing pre-S(120-145) in a selected patient is shown in 9 Fig. 13. 10 Anti-pre-S protein antibodies are detected in 11 human sera early during acute hepatitis type B. 12 antibodies recognizing the peptides were detected during 13 HBsAg antigenemia before antibodies to the S-protein 14 (anti-HBs) or to hepatitis B core antigen (anti-HBc) were 15 detectable. After development of the latter two antibodies, 16 the level of antibodies with anti-pre-S specificity 17 declined. Variations of this pattern of anti-pre-S 18 development among patients with hepatitis B were observed. 19 In some cases, antibodies recognizing the synthetic peptides 20 were present even before HBsAg was detected in plasma, or 21 when HBsAg never appeared in blood and the only marker for 22 hepatitis B was anti-HBc and later anti-HBs. 23 Antibodies to pre-S(120-145) were measured by RIA. 24 Similar results were obtained by assaying antibodies to 25 pre-S(12-32). HBsAg, anti-HBs and antibodies to hepatitis B 26 core antigen (anti-HBc) were assayed using commercial test 27 kits (Abbot Laboratories, North Chicago, Illinois). 28 broken line at the end of bars corresponding to the 29

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different markers of HBV infection indicates positivity at

1 the termination of surveilance. Antibody titers represent 2 the highest dilution of serum at which radioactivity counts 3 corresponding to the specimens divided by counts corresponding to equally diluted control serum were > 2.1. 5 Humans vaccinated with pepsin-treated HBsAg 6 (Hilleman, M.R., Buynak, E.B., McAleer, W.J., McLean, A.A., 7 Provost, P.J. & Tytell, A.A. in Viral Hepatitis, 1981 8 International Sympsosium (eds. Szmuness, W., Alter, H.J. & 9 Maynard, J.E.) 385-397 (Franklin Institute Press, 10 Philadelphia, PA, 1982)), (pepsin treatment removes all 11 anti-pre-S(120-145) reactive material), or with HBsAg 12 produced in yeast (devoid of pre-S gene coded sequences 13 (McAleer, W.J. Buynak, E.B. Maigetter, R.Z., Wambler, D.E., 14 Miller, W.J., Hillemann, M.R. Nature, (London), 307, 178-180 15 (1984); did not develop detectable antibodies recognizing 16 either of the two synthetic peptides. On the other hand, 7 17 out of 12 individuals who received a vaccine consisting of 18 intact HBsAg (McAuliffe, V.J., Purcell, R.H., Gerin, J.L. & 19 Tyeryar, F.J. in Viral Hepatitis (eds Szmuness, W., Alter, 20 H.J. & Maynard, J.E.) 425-435, Franklin Institute Press, 21 Philadelphia, PA) developed these antibodies. 22 individuals also had the highest antibodiy response to the 23 S-protein, as measured by the AUSAB test (Abbott), 24 suggesting that a lack of detectable response to the pre-S 25 protein was due to the sensitivity limits of the test. 26 this respect, it is of importance that the hepatitis B 27 vaccine heretofore used, the production of which involves 28 pepsin treatment of HBCAg, although highly efficient in 29 apparently healthy individuals, has had low immunogencity 30

and no protective effect in hemodialysis patients (Stevens, C.E., Alter, H.J., Taylor. P.E., Zang, E.A., Harley, E.J. & Szmuness, W., N. Engl. J. Med., 311, 496-501 (1984)). Other vaccines produced without pepsin treatment do not seem to have this defect (Desmyter, J. in <u>Viral Hepatitis and Liver Disease</u> (eds Vyas, G.N., Dienstag, J.L. & Hoofnagle, J.), in press Grune and Stratton, Orlando, Fl. 1984).

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#### Example 25

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# RIA Tests of Preparations Containing HBV-specific proteins

Antibodies to the S-protein were removed from rabbit anti-serum against HBV particles by affinity chromatography (Neurath, A.R., Trepo, C., Chen, M., Prince, A.M., J. Gen. Virol., 30, 277-285 (1976) - See Fig. 14. The tested antigens were: HBV particles and tubular forms (e, A); approximately 20 nm spherical particles of HBsAg isolated from plasma  $(0, \Delta)$ ; and the latter particles treated with pepsin (1 mg/ml HBsAg, 50ug/ml pepsin in 0.1 M glycine-HCl, pH 2.2, 2 hours at 37°C) (□). The RIA tests were performed as described in Neurath; A.R., Kent, S.B.H., Strick, N., Science, 224, 392-395 (1984). The concentration of HBsAg S-protein was adjusted to the same level in all preparations tested as based on RIA tests (AUSRIA, Abbot Laboratories). HBV particles (contaminated with tubular forms of HBsAq) were concentrated from serum approximately 100x by centrifugation for 4 hours at 25,000 rpm in a Spinco 35 rotor. The concentrate (2 ml) was layered over a discontinuous gradient consisting of 11 ml of each 20, 10

and 5% sucrose (w/w) in 0.14 M NuCl-0.01 M Tris-0.02% NaN3,

pH 7.2 (TS) and centrifuged for 16 hours at 25,000 rpm in a

Spinco rotor SW 27. The final pellet was resuspended in TS.

HBV particles were recognized much more

efficiently than purified approximately 22 nm spherical

efficiently than purified approximately 22 nm spherical particles in RIA tests based on polystyrene beads coated with either anti-pre-S(120-145) or with rabbit antibodies to HBV particles. Treatment of HBsAg with pepsin, a step used in preparing some current hepatitis B vaccines, resulted in an approximately 10<sup>3</sup>-fold decrease in reactivity with anti-pre-S(120-145). HBsAg from vaccines derived either from infected plasma (Hilleman, M.R., et al, 1982) <a href="mailto:supra">supra</a>), or produced in yeast McAleer et al (1984), <a href="mailto:supra">supra</a>), had <a href="mailto:supra">had</a>

In reverse tests, beads coated with HBsAg, with HBV particles, with pepsin-treated HBsAg, or with HBsAg corresponding to the vaccines mentioned above were utilized. IgG antibodies (from different rabbit antisera to pre-S sequences) reacting with the beads were assayed based on the subsequent attachment of labeled anti-rabbit IgG. Positive results using anti-pre-S(120-145) were obtained only with beads coated with intact HBsAg or with HBV particles. Anti-pre-S(12-32) reacted exclusively with HBV-coated beads.

# Example 26 Involvement of pre-S Gene Coded HBV Domains In

### Attachment to Cell Receptors

It has been suggested that the 55 C-terminal amino acids of the pre-S protein mediate the attachment of HBsAg to human albumin polymerized by glutaraldehyde (pHSA) and

that this attachment plays an essential role in the in vivo adsorption of HBV to hepatocytes (Machida, A. et al, Gastroenterology, 86, 910-918, (1984); Machida, A. et al, Gastroenterology, 85, 268-274, (1983). However, there is no compelling evidence to support the role of the pHSA-HBV interaction in infection of liver cells by HBV. In addition, both HBsAg containing or lacking these 55 amino acid residues react with pHSA (Fig. 15), albeit the reaction is enhanced by the presence of the pre-S gene coded sequences. The RIA tests involved in Fig. 15 were conducted as described in Neurath, A.R., Strick, N. Intervirology, 11, 128-132 (1979).

To explore directly the reaction of HBsAg with liver cells, an assay system based on the attachment of liver cells to insolubilized HBsAg was developed.

HBsAg (HBV) was attached to

N-N'-p-phenylenedimaleimide-derivatized sulfhydryl cellulose under conditions described for linking of pre-S(120-145), as described above. About 4 mg of HBsAg was linked to 1 g of the cellulose derivative. A control cellulose derivative was prepared by linking bovine serum albumin to the activated matrix. Forty mg of the cellulose derivative suspended in TS containing 10 mg/ml of bovine serum albumin (TS-BSA) were mixed with approximately 2 x 10<sup>6</sup> washed Hep G2 human hepatoma cells (see Aden, D.P., Fogel, A., Plotkin, S., Damjanov, J., Knowles, B.B., Nature (London), 282, 615-617 (1979) suspended in TS-BSA and incubated for 30 min at 37°C, followed by 1 hour at 4°C. HeLa cells and Clone 9 normal rat liver cells (American Type Culture Collection)

were used as controls. The cell-cellulose mixtures were layered on top of 1 ml cf 33% (w/w) Hypaque and centrifuged for 2 minutes at 3,000 rpm. The cellulose derivative with attached cells pelleted under these conditions. Unattached cells recovered from the Hypaque-TS-BSA interphase were diluted 5-fold in TS-BSA and pelleted by centrifugation. The relative proportion of adsorbed and unadsorbed cells was 8 determined by measurement of lactate dehydrogenase (LDH) 9 activity in appropriate aliquots of cell lysates obtained 10 after exposure to the detergent Triton X-100 (5 mg/ml in 11 H2O). LDH activity was determined using diagnostic kit No. 12 500 (Sigma). 13 Approximately 80 to 95% of human hepatoma Hep G2 14 cells (Aden, D.P. supra) attached to immobilized HBsAg in 15 this assay. The attachment of control cells (HeLa, rat 16 hepatocytes) was in the range of 10 to 20%. About 10% of 17 Hep G2 cells attached to control cellulose. In the presence 18 of anti-pre-S(120-145) and anti-pre-S(12-32) IgG (15 mg/ml), 19 the adsorption of Hep G2 cells to HBsAg-cellulose decreased 20 to 60 and 30%, respectively. A mixture of both antibodies 21 (7.5 mg/ml of IgG each) caused a decrease of cell adsorption 22 to 20%, indistinguishable from background levels. 23 Normal rabbit IgG, as well as antibodies to the 24 S-protein (elicited by immunization with pepsin-treated 25 HBsAg), failed to diminish the cell attachment, despite high

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It will be appreciated that the instant specification and claims are set forth by way of

dilution in the AUSAB test).

levels of anti-HBs present in this serum (positive at a  $10^{-6}$ 

1	ill	ustrati	on an	d not	lim	itat	ion	and	that	vari	ous	
2	-										departing	from
3	the	spirit	and	scope	of t	the	pres	ent	inve	ntion		
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-	WHAT IS CLAIMED IS:
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3 `	1. A hepatitis B peptide immunogen comprising
4	peptide containing an amino acid chain
5	corresponding to at least six consecutive amino acids within
6	the pre-S gene coded region of the envelope of HBV, said
7	peptide immunogen free of an amino acid sequence
8	corresponding to the naturally occurring envelope proteins
9	of hepatitis B virus.
10	<ol> <li>A hepatitis B peptide immunogen according to</li> </ol>
11	claim 1, wherein said chain of amino acids is between
12	sequence position pre-S 120 and pre-S 174.
13	3. A hepatitis B peptide immunogen according to
14	claim 2, wherein said chain of amino acids includes
15	N-terminal methionine at sequence position pre-S 120.
16	4. A heptatis B peptide immunogen according to
17	claim 1, wherein said chain of amino acids is between
18	sequence position pre-S 1 and pre-S 120.
19	5. A hepatitis B peptide immunogen according to
20	claim 1, wherein said peptide contains a chain of at least
21	10 amino acids.
22	6. A hepatitis B peptide immunogen according to
23	claim 1, wherein said peptide contains a chain of at least
24	15 amino acids.
25	7. A hepatitis B peptide immunogen according to
26	claim 1, wherein said peptide contains a chain of at least
27	20 amino acids.
28	20 dillio dolds.
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ı	8. A hepatitis B peptide immunogen according to
2	claim 1, wherein said peptide contains a chain of at least
3	26 amino acids.
4	<ol> <li>A hepatitis B peptide immunogen according to</li> </ol>
5	claim 8, wherein said chain is between and including
6	sequence positions pre-S 120 and pre-S 174.
7	10. A hepatitis B peptide immunogen according to
8	claim 9, wherein said chain includes N-terminal methionine
9	at sequence position pre-S 120.
10	11. A hepatitis B peptide immunogen according to
11	claim 1, wherein said chain is between and including
12	sequence position pre-S 15 and pre-S 120.
13	12. A hepatitis B peptide immunogen according to
14	claim 1, wherein said chain is between and including
15	sequence position pre-S 15 and pre-S 55.
16	13. A hepatitis B peptide immunogen according to
17	claim 1, wherein said chain is between and including
18	sequence position pre-S 90 and pre-S 120.
19	14. A hepatitis B peptide immunogen according to
20	claim 1, wherein said chain is between and including
21	sequence position pre-S 10 and pre-S 40.
22	15. A hepatitis B peptide immunogen according to
23	claim 1, wherein said chain corresponds to amino acids in
24	the ayw subtype of the pre-S region.
25	16. A hepatitis B peptide immunogen according to
26	claim 1, wherein said chain corresponds to amino acids in
27	the adyw subtype of the pre-S region.
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1	17. A hepatitis B peptide immunogen according to
2	claim 1, wherein said chain corresponds to amino acids in
3	the adw2 subtype of the pre-S region.
4	18. A hepatitis B pept.le immunogen according to
5	claim 1, wherein said chain corresponds to amino acid in the
6	adw subtype of the pre-S region.
7	19. A hepatitis B peptide immunogen according to
8	claim 1, wherein said chain corresponds to amino acids in
9	the adr subtype of the pre-S region.
10	20. A hepatitis B peptide immunogen according to
11	claim 1, wherein said peptide immunogen is free of any serum
12	proteins.
13	
14	21. A heptatis B peptide immunogen according to
15	claim 1, wherein said chain is MQWNSTAFHQTLQDPRVRGLYLPAGG.
16	22. A hepatitis B peptide immunogen according to
17	claim 1, wherein said chain is MGTNLSVPNPLGFFPDHQLDP.
18	23. A hepatitis B peptide immunogen according to
19	claim 1, wherein said chain is PAFGANSNNPDWFNPVKDDWP.
20	24. A hepatitis B peptide immunogen according to
21	claim 1, wherein said chain is PQAMQWNSTAFHQTLQDP.
22	25. A hepatitis B peptide immunogen according to
23	claim 1, wherein said chain is PASTNRQSGRQPTPISPPLRDSHP.
24	26. A hepatitis B peptide immunogen according to
25	claim 1, wherein said chain is PAPNIASHISSISARTGDP.
26	27. A hepatitis B peptide immunogen according to
27	claim 1, wherein said chain is MGGWSSKPRKGMGTNLSVPNP.
28	28. A hepatitis B peptide immunogen according to
29	claim 1, wherein said chain is PAFGANSNNPDWDFNPVKDDWP.
- /	

1	. 29. A hepatitis B peptide immunogen according to
2	claim 1, wherein said chain is QVGVGAFGPRLTPPHGG.
3	30. A hepatitis B peptide immunogen according to
4	claim 1, wherein said chain is MGGWSSKPRKG.
5	31. A hepatitis B peptide immunogen according to
6	claim 1, wherein said chain is MGGWSSKPRQG.
7	32. A hepatitis B peptide immunogen according to
8	claim 1, wherein said peptide immunogen is free of an amino
9	acid sequence corresponding to the entire S gene coded
10	region of the env gene product of hepatitis B virus.
11	33. A hepatitis B peptide immunogen according to
12	claim 1, wherein said peptide has no more than 100 amino
13	acids.
14	34. A hepatitis B peptide immunogen according to
15	claim 1, wherein said peptide has no more than 40 amino
16	acids.
17	35. A hepatitis B peptide immunogen according to
18	claim 1, wherein said peptide has no more than 30 amino
19	acids.
20	36. A hepatitis B peptide immunogen according to
21	claim 1, wherein said peptide is capable of forming
22	neutralizing antibodies to hepatitis B virus in a humans.
23	37. A hepatitis B peptide immunogen according to
24	claim 1, wherein said peptide is linked to a carrier.
25	38. A hepatitis B peptide immunogen according to
26	claim 37, wherein said peptide is covalently linked to a
27	carrier.
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_	39. A hepatitis B peptide immunogen according to
2	claim 38, wherein said peptide is covalently linked to a
3	lipid vesicle carrier.
4	40. A hepatitis B peptide immunogen according to
5	claim 39, wherein said lipid vesicle is stabilized by
6	cross-linking.
7	41. A hepatitis B peptide immunogen according to
8	claim 1, wherein said chain is pre-S(120-145).
9	42. A hepatitis B peptide immunogen according to
10	claim 1, wherein said chain is pre-S(12-32).
11	43. A hepatitis B peptide immunogen according to
12	claim 1, wherein said chain is pre-S(117-134).
13	44. A hepatitis B peptide immunogen according to
14	claim 1, wherein said chain is pre-S(94-117).
15	45. A hepatitis B peptide immunogen according to
16	claim 1, wherein said chain is pre-S(153-171).
17	
18	to the property of the propert
19	claim 1, wherein said chain is pre-S(1-21).
20	47. A hepatitis B peptide immunogen according to
21	claim 1, wherein said chain is pre-S(57-73).
22	48. A carrier for a peptide comprising a lipid
23	vesicle stabilized by cross-linking and having covalently
24	bonded active sites on the outer surface thereof to bind the
25	peptide to the outer surface of the carrier.
26	49. A carrier according to claim 48, wherein said
27	active sites are selected from the group consisting of
28	-COOH, -CHO, -NH <sub>2</sub> and -SH.
29	50. A carrier according to claim 49, wherein said
30	lipid vesicle contains an amino moiety selected from the

1	group consisting of aminoalkane, diaminoalkane, aminoalkene
2	and diaminoalkene having 10 to 18 carbon atoms and said
3	lipid vesicle is stabilized by contacting said lipid vesicle
4	with a polyaldehyde.
5	51. A carrier according to claim 50, wherein said
6	polyaldehyde is a bifunctional aldehyde.
7	52. A carrier according to claim 51, wherein said
8	bisaldehyde is glutaraldehyde.
9	53. A carrier according to claim 50, wherein said
10	amino moiety is stearylamine.
11	54. A carrier according to claim 52, wherein said
12	amino moiety is stearylamine.
13	55. A carrier according to claim 48, wherein said
14	lipid vesicle contains fatty acids having 12 to 18 carbon
15	atoms and said lipid vesicle is stabilized with a
16	carbodiimide.
17	56. A carrier according to claim 55, wherein said
18	fatty acid is stearic acid and said carbodiimide is
19	N-ethyl-N'(dimethyl-aminopropyl)-carbodiimide.
20	57. A carrier according to claim 48, wherein said
21	lipid vesicle contains fatty acid aldehyde.
22	58. A peptide linked to a carrier comprising a
23	peptide having -SH groups and a carrier comprising a liquid
24	vesicle containing an amino moiety selected from the group
25	consisting of aminoalkane, diaminoalkane, aminoalkene,
26	diaminoalkene having 10 to 18 carbon atoms activated by a
27	polyaldehyde and further activated by cysteine.
28	59. A peptide linked to a carrier according to
29	5). A popular limbur of a dalile door and

claim 58, wherein said -SH groups are supplied by cysteine.

-	60. A peptide linked to a carrier comprising a
2	peptide having -SH groups and a carrier comprising a lipid
3 `	vesicle containing fatty acid mercaptan.
4	61. A peptide linked to a carrier according to
5	claim 60, wherein said fatty acid mercaptan is
6	octadecanethiol.
7	62. A peptide linked to a carrier according to
8	claim 60, wherein said fatty acid mercaptan contains lipid
9	vesicle activated with a dimaleimide.
10	63. A peptide linked to a carrier according to
11	claim 52, wherein said dimaleimide is N-N'-phenylanedi-
12	maleimide.
13	64. A peptide linked to a carrier comprising a
14	peptide activated by a carbodiimide and a carrier comprising
15	a lipid vesicle containing an amino moiety selected from the
16	group consisting of aminoalkane, diaminoalkane, aminoalkene,
17	diaminoalkene having 10 to 18 carbon atoms.
18	65. A peptide linked to a carrier according to
19 20	claim 64, wherein said amino moiety is stearylamine.
21	66. A peptide linked to a carrier according to
22	claim 64, wherein said carbodiimide is
23	N-ethyl-N' (dimethylaminopropyl) -carbodiimide.
24	67. A peptide linked to a carrier comprising a
25	peptide activated by a carbodiimide and a carrier comprising
26	a lipid vesicle stabilized by a polyaldehyde and further
27	derivatized by reaction with a water-soluble diaminoalkane.
28	68. A peptide linked to a carrier according to
29	claim 67, wherein said carbodiimide is N-ethyl-N'(di-

1 methylaminopropyl)-carbodiimide, said polyaldehyde is 2 glucaraldehyde and said diaminoalane is ethylenediamine. 3 A method of forming a carrier comprising 69. contacting a lipid vesicle containing an amino moiety 5 selected from the group consisting of aminoalkane, 6 diaminoalkane, aminoalkene and diaminoalkene having 10 to 18 carbon atoms with a polyaldehye. 8 70. A method according to claim 69, wherein said 9 amino moiety is stearylamine. 10 A method according to claim 69, wherein said 71. 11 polyaldehyde is glutaraldehye. 12 A method of forming a carrier comprising 13 contacting a lipid vésicle containing fatty acid having 12 14 to 18 carbon atoms with a carbodiimide. 15 73. A method of forming a carrier according to 16 claim 72, wherein said fatty acid is stearic acid and said 17 carbodiimide is N-ethyl-N'(dimethylaminopropyl)-18 carbodiimide. 19 A method of forming a carrier comprising 20 contacting a lipid vesicle with a fatty acid aldehyde. 21 75. A method of linking a peptide to a carrier 22 comprising contacting a peptide having -SH groups with a 23 carrier comprising a lipid vesicle containing an amino 24 moiety selected from the group consisting of aminoalkane, 25 diaminoalkane, aminoalkene and diaminoalkene having 10 to 18 26 carbon atoms and contacting said lipid vesicle with a

comprising contacting a peptide having -SH groups with a

A method of linking a peptide to a carrier

polyaldehyde and cysteine.

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1	carrier comprising a lipid vesicle having fatty acid
2	mercaptan.
3	77. A method of linking a peptide to a carrier
4	according to claim 76, wherein said fatty acid mercaptan is
5	octadecanediol.
6	78. A method of linking a peptide to a carrier
7	according to claim 77, wherein said fatty acid mercaptan
8	containing lipid vesicle is contacted with a dimaleiimide.
9	79. A method according to claim 78, wherein said
10	dimaleiimide is N-N'-phenylanedimaleimide.
11	80. A method of linking a peptide to a carrier
12	comprising contacting a peptide, said peptide contacted with
13	
14	a carbodiimide, with a carrier, said carrier comprising an
15	lipid vesicle containing an amino moiety selected from the
16	group consisting of aminoalkane, diaminoalkane, aminoalkene
17	and diaminoalkene having 10 to 18 carbon atoms.
18	81. A method according to claim 80, wherein said
19	amino moiety is stearylamine and said carbodiimide is
20	N-ethyl-N'(dimethylaminopropyl)-carbodiimide.
21	82. A method of linking a peptide to a carrier
22	comprising contacting a peptide activated by a carbodiimide
23	and a carrier comprising a lipid vesicle stabilizied by a
24	polyaldehyde and further reacted with a water-soluble
25	diaminoalkane.
26	83. A method of linking a peptide to a carrier
27	according to claim 82, wherein said polyaldehyde is
28	glutaraldehyde and said diaminoalkane is ethylenediamine.
29	84. A peptide comprising an amino acid chain
30	corresponding to at least six consecutive amino acids within

*	the pre-S gene coded region or the envelope of HBV, said
2	peptide free of an amino acid sequence corresponding to the
3	naturally occurring envelope proteins of hepatitis B virus.
4	85. A peptide according to claim 84, wherein
5	said chain of amino acids is between sequence position pre-S
6	120 and pre-S 174.
7	86. A peptide according to claim 85, wherein
8	said chain of amino acids includes N-terminal methionine at
9	sequence position pre-S 120.
LO	87. A peptide according to claim 84, wherein
Ll	said chain of amino acids is between sequence position pre-s
L2	1 and pre-S 120.
13	88. A peptide according to claim 84, wherein
L 4	said peptide contains a chain of at least 10 amino acids.
L5	89. A peptide according to claim 84, wherein
L6	
L7	said peptide contains a chain of at least 15 amino acids.
18	90. A peptide according to claim 84, wherein
L9	said peptide contains a chain of at least 20 amino acids.
20	91. A peptide according to claim 84, wherein
21	said peptide contains a chain of at least 26 amino acids.
22	92. A peptide according to claim 91, wherein
23	said chain is between and including sequence positions pre-S
24	120 and pre-S 174.
25	93. A peptide according to claim 92, wherein
26	said chain includes N-terminal methionine at sequence
27	position pre-S 120.
28	94. A peptide according to claim 84, wherein
29	said chain is between and including sequence position pre-S

15 and pre-S 120.

1	95. A peptide according to claim 84, wherein
2	said chain is between and including sequence position pre-S
3	15 and pre-S 55.
4	96. A peptide according to claim 84, wherein
5	said chain is between and including sequence position pre-S
6	90 and pre-S 120.
7	97. A peptide according to claim 84, wherein
8	said chain is between and including sequence position pre-S
9	10 and pre-S 40.
10	98. A peptide according to claim 84, wherein
11	said chain corresponds to amino acids in the ayw subtype of
12	the pre-S region.
13	99. A peptide according to claim 84, wherein
14	said chain corresponds to amino acids in the adyw subtype of
15	the pre-S region.
16	100. A peptide according to claim 84, wherein
17	said chain corresponds to amino acids in the adw2 subtype of
18	the pre-S region.
19	101. A peptide according to claim 84, wherein
20	said claim corresponds to amino acids in the adw subtype of
21	the pre-S region.
22	102. A peptide according to claim 84, wherein
23	said chain corresponds to amino acids in the adr subtype of
24	the pre-S region.
25	103. A peptide according to claim 84, wherein
26	said peptide is MQWNSTAFHQTLQDPRVRGLYLPAGG.
27	104. A peptide according to claim 84, wherein
28	said peptide is MGTNLSVPNPLGFFPDHQLDP.
29	I -!

1	•	10	)5.	A	peptide	according	to	claim	84,	wherein	
2	said	peptide	is	P	AFGANSNPI	OWFNPVKDDW	P.				
3		10	6.	A	peptide	according	to	claim	84,	wherein	
4	said	peptide	is	Pζ	QAMQWNST <i>i</i>	AFHQTLQDP.					
5		. 10	7.	A	peptide	according	to	claim	84,	wherein	
6	said	peptide	is	P	ASTNRQSGI	RQPTPISPPL	RDSI	HP.			
7		10	8.	A	peptide	according	to	claim	84,	wherein	
8	said	peptide	e is	PZ	APNIASHIS	SSISARTGDP	•				
9		10	9.	A	peptide	according	to	claim	84,	wherein	
LO	said	peptide	e is	MO	GWSSKPRI	KGMGTNLSVP	NP.				
11		11	.0.	A	peptide	according	to	claim	84,	wherein	
L2	said	peptide	e is	PI	AFGANSNNI	PDWDFNPVKD	DWP.	•			
L3					_	according			84,	wherein	
14	said					RLTPPHGG.			-		
15						according	to	claim	84,	wherein	
16	said				GWSSKPRI	_			·		
17						according	to	claim	84.	wherein	
18	said				GWSSKPRI				•		
19						according	to	claim	84.	wherein	
20	saið					n amino ac					ling
21						d region o		_		_	_
22		titis B		_		<b>.</b>					-
23	nopu					according	to	claim	84.	wherein	
24	caid					than 100 a			•	***************************************	
25	3414					according				whorein	
26	anid					than 40 am				MIGLETII	
27	Salu										
28						according				wnerein	
29	said	peptid	e na	s :	no more	than 30 am	ııno	acids	•		

1	118. A peptide according to claim 84, wherein
2	said chain is pre-S(120-145).
3	119. A peptide according to claim 84, wherein
4	said chain is pre-S(12-32).
5	120. A peptide according to claim 84, wherein
6	said chain is pre-S(117-134).
7	121. A peptide according to claim 84, wherein said
8	chain is pre-S(94-117).
9	122. A peptide according to claim 84, wherein
10	said chain is pre-S(153-171).
11	123. A peptide according to claim 84, wherein
12	said chain is pre-S(1-21).
13	124. A peptide according to claim 84, wherein
14	said chain is pre-S(57-73).
15	125. A process for the detection of antigens
16	coded for the pre-S gene in sera of HBV infected animals
17	comprising:
18	(a) coating a solid substrate with
19	antibodies to a peptide having an amino acid chain
20	corresponding to at least six consecutive amino acids within
21	the pre-S gene coded region of the envelope of HBV, said
22	peptide free of an amino acid sequence corresponding to the
23	naturally occurring proteins of HBV;
24	(b) washing the coated substrate;
25 '	(c) contacting the washed coated substrate
26	with a protein-containing solution;
27	(d) washing the substrate from step c;
28	
29	
30	with a sample suspected to contain HBV or HBsAg;

-	(f) washing the substrate from step e;
2	(g) adding radiolabeled antibody, said
3	antibody being an antibody to the peptide or HBsAg, to the
4	substrate from step f;
5	(h) incubating the substrate from step g;
6	(i) washing the substrate from step h; and
7	(j) subjecting the substrate of step i to
8	counting in a gamma counter.
9	126. A process for the detection of antigen coded
10	for by the pre-S gene in sera of HBV infected animals
11	according to claim 125, wherein said substrate is
12	polystyrene beads.
13	127. A process for the detection of antigen coded
14	for by the pre-S gene in sera of HBV infected animals
15	according to claim 125, wherein said protein-containing
16	solution contains bovine serum albumin or gelatin.
17	128. A process for the detection of antigens coded
18	for the pre-S gene in sera of HBV infected animals
19	comprising:
20 21	(a) coating a solid substrate with
22	antibodies to a peptide having an amino acid chain
23	corresponding to at least six consecutive amino acids within
24	the pre-S gene coded region of the envelope of HBV, said
25	peptide free of an amino acid sequence corresponding to the
26	naturally occurring proteins of HBV;
27	(b) washing the coated substrate;
28	(c) contacting the washed coated substrate
29	with a protein-containing solution;
30	(d) washing the substrate from step c:

1	(e) incubating the substrate from step d
2 .	with a sample suspected to contain HBV or HBsAg;
3	(f) washing the substrate from step e;
4	(g) adding enzyme labeled antibody, said
5	antibody being an antibody to the peptide or HBsAg, to the
6	substrate from step f;
7	(h) incubating the substrate from step g;
8	(i) washing the substrate from step h;
9	(j) subjecting the substrate of step i to
10	ELISA and comparing the results of said ELISA to ELISA
11	results from normal sera utilized as a control.
12	129. A process for the detection of antigens coded
13	for by the pre-S gene in sera of HBV infected animals
14	according to claim 128, wherein said substrate is
15	polystyrene beads.
16	130. A process for the detection of antibodies to
17	the pre-S region of hepatitis B virus comprising:
18	(a) adsorbing on a solid substrate,
19	containing binding sites thereon, a peptide having an amino
20	acid chain corresponding to at least six consecutive amino
21	acids within the pre-S gene coded region of the envelope of
23	HBV, said peptide free of an amino acid sequence
24	corresponding to the naturally occurring envelope proteins
25	of hepatitis B virus,
26	(b) contacting the substrate from step a
27	with a material to saturate the binding sites thereon,
28	(c) washing the substrate from step b,
29	
30	

1 .	(d) contacting the substrate from step c
2	with a specimen comprising human sera to form a first
3	resultant mass,
4	(e) incubating the resultant mass of step d,
5	(f) washing the resultant mass of step e,
6	(g) adding radiolabeled antibodies to human
7	IGG or IGM to the resultant mass of step f to form a second
8	resultant mass,
9	(h) subjecting the second resultant mass of
10	step g to counting in a gamma counter,
11	(i) subjecting normal sera utilized as a
12	control to steps a to h and
13	(j) comparing the counts of steps h and i.
14	131. A process according to claim 130, wherein
15	said substrate is polystyrene beads.
16	132. A process for the detection of antibodies to
17	the pre-S region of hepatitis B virus comprising:
18	(a) adsorbing on a solid substrate,
19	containing binding sites thereon, a peptide having an amino
20	acid chain corresponding to at least six consecutive amino
21	acids within the pre-S gene coded region of the envelope of
22	HBV, said peptide free of an amino acid sequence
23	corresponding to the naturally occurring envelope proteins
24	of hepatitis B virus,
25	(b) contacting the substrate from step a
26	
27	with a material to saturate the binding sites thereon,
28	(c) washing the substrate from step b,
29	

1	(d)	contacting the substrate from s ? c
2	with a specimen co	mprising human sera to form a first
3 ,	resultant mass,	
4	(e)	incubating the resultant mass of step d,
5	(f)	washing the resultant mass of step e,
6	(g)	adding enzyme labeled antibodies to
7	human IgG or IgM t	o the resultant mass of step f to form a
8	second resultant m	
9	(h)	subjecting the second resultant mass of
10	step g to ELISA,	
11	-	subjecting a normal sera utilized as a
12	control to ELISA,	
13	(j)	•
14	and i.	-
15		process according to claim 132, wherein
16	-	polystyrene beads.
17		liagnostic test kit for detecting antigens
18		-S gene of HBV in a test sample, comprising
19	a.	a solid substrate containing protein
20		reon, said substrate coated with antibodies
21		ng an amino acid chain corresponding to at
22		tive amino acids within the pre-S gene
23		
24		he envelope of HBV, said peptide free of an
25	_	ce corresponding to the naturally occurring
26	proteins of HBV,	
27		a protein-containing solution to
28		binding sites on the solid substrate, and
29	c.	
3.0	labelled antibody	, said antibody to either the peptide or

1	HBsAg.
2	135. A diagnostic test kit for detecting
3	antibodies to the pre-S region of hepatitis B virus in a
4	test sample, comprising
5	a. a solid substrate containing protein
6	binding sites thereon, said substrate having adsorbed
7	thereon a peptide having an amino acid chain corresponding
8	to at least six consecutive amino acids within the pre-S
9	gene coded region of the envelope of HBV, said peptide free
10	of an amino acid sequence corresponding to the naturally
11	occurring proteins of HBV, the solid substrate exposed to
12	a protein-containing solution to saturate protein binding
13	sites on the solid substrate, and
14	b. a given amount of radiolabeled or enzyme
15	labelled antibodies to human IgG or IgM.
16	136. A process for detecting antibodies to the
17	pre-S region of hepatitis B virus in a sample which
18	comprises:
20	<ul> <li>a) contacting the sample with a solid substrate</li> </ul>
21	coated with a non-labelled peptide containing an amino acid
22	chain corresponding to at least six consecutive amino acids
23	within the pre-S gene coded region of the enveloping of HBV,
24	the peptide free of an amino acid sequence corresponding to
25	the naturally occurring envelope proteins of hepatitis B
26	virus, incubating and washing said contacted sample;
27	b) contacting the incubated washed product
28	obtained from step a above with a labelled peptide
29	containing an amino acid chain corresponding to at least six
30	consecutive amino acids within the pre-S gene coded region

of the envelope of HBV, said peptide free of an amino acid 2 sequence corresponding to the naturally occurring envelope 3 protein of hepatitis B virus, incubating and washing the 4 resultant mass; and 5 determing the extent of labelled peptide c) 6 present in the resultant mass obtained by step b above. 7 137. A process according to claim 136, wherein 8 a solid substrate is rendered substantially free of 9 available protein binding sites. 10 138. A process according to claim 137, wherein 11 the solid substrate is contacted with a protein binding site 12 occupier. 13 139. A process according to claim 139, wherein 14 the occupier is albumin. 15 A process for detecting antibodies to the 140. 16 pre-S region of hepatitis B virus in a sample comprising: 17 contacting the sample with a solid substrate 18 coated with a non-labelled peptide containing an amino acid 19 chain corresponding to at least six consecutive amino acids 20 within the pre-S gene coded region of the envelope of HBV, 21 the peptide free of an amino acid sequence corresponding to 22 the naturally occurring envelope proteins of hepatitis B 23 virus, incubating and washing said contacted sample; 24 contacting the incubated washed product 25 obtained from step a above with labelled antibody to human 26 or animal immunoglobulin product by contact with an 27 immunogen comprising a peptide corresponding to at least six 28 consecutive amino acids within the pre-S gene coded region 29

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of the envelope of HBV, said peptide immunogen free of an

•	amino acid sequence corresponding to the naturally occurring
2	envelope proteins of hepatitis B virus, incubacing and
3	washing the contacted sample, and
4	c) determining the exten of labelled antibody
5	present in the resultant mass of step b.
6	141. A process for detecting HBV or HBsAg in a
7	sample comprising:
8	a) contacting a first portion of a composition
9	containing an antibody produced by introducing into an
10	animal or human an immunogen comprising a peptide
11	corresponding to at least six consecutive amino acids within
L2	the pre-S gene coded region of the envelope of HBV, said
13	peptide immunogen free of an amino acid sequence
14	corresponding to the naturally occurring envelope proteins
15, 16	of hepatitis B virus with a mixture of said sample and said
L7	immunogen which has been labelled, incubating and washing
18	said first protein;
19	b) contacting a second portion of said
20	composition containing antibody with the same amount of said
21	labelled immunogen in an antigen free control, incubating
22	and washing said second portion;
23	c) adding the same amount of Staphylococci
24	bearing protein A to each of the compositions of steps a and
25	b above, incubating both of said compositions, centrifuging
26	each of said compositions and separating liquid from the
27	solids therein;
28	d) determining the extent of labelled immunogen
29	in each of the resultant compositions from step c above, and
20	a) companies the malating amount of labelled

-	immunogen in each such that if the activity of the resultant
2	composition containing the first portion is less than the
3 .	activity for the resultant composition of the second
4	portion, then the sample contains HBV or HBsAg.
5	142. A diagnostic test kit for detecting
6	hepatitis B virus in sera comprising
7	a) a given amount of antibody to a peptide
8	containing an amino acid chain corresponding to at least six
9	consecutive amino acids within the pre-S gene coded region
10	of the envelope of HBV, said peptide being free of an amino
11	acid chain corresponding to the naturally occurring envelope
12	proteins of hepatitis B virus, the antibody being bound to a
13	solid support,
14	b) labelled antibody to the peptide or to
15	hepatitis B virus.
16	143. A diagnostic test kit for detecting
17	hepatitis B virus in sera according to claim 142, which
18	further comprises a set of instructions for effecting an
19	immunoassay wherein the effect of formation of an immune
20	complex is revealed by said labelled antibody.
21	144. A diagnostic test kit for detecting
22	
23	hepatitis B virus in sera according to claim 143, wherein said antibody is insolubilized on a water insoluble solid
24	
25	support.
26	145. A diagnostic kit for detecting the presence
27	of antibodies to hepatitis B virus comprising
28	a) a given amount of a peptide containing
29	an amino acid chain corresponding to at least six
30	consecutive amino acids within the pre-S gene coded region

of the envelope of HBV, said peptide being free of an amino

acid chain corresponding to the naturally occurring envelope

proteins of hepatitis B virus,

b) labelled antibodies to human IgG or IgM.

of antibodies to hepatitis B virus according to claim 145, which further comprises a set of instructions for effecting an immunoassay, wherein the extent of formation of an immune complex is revealed by said labelled antibodies.

147. A diagnostic test kit for detecting hepatitis B virus in sera according to claim 145, wherein said peptide is insolubilized upon a water insoluble solid support.

148. A process for predicting the outcome of hepatitis B infection which comprises carrying out an immunoassay on serum of a human to detect the presence of an antibody to an antigen coded for by the pre-S gene coded region of the envelope of hepatitis B virus employing the peptide immunogen of claim 1 at regular intervals and evaluating the data.

been vaccinated with a vaccine against hepatitis B has become immune to hepatitis B virus which comprises effecting a plurality of immunoassays of serum from such human to determine if there are antibodies in said serum to an antigen coded by the pre-S gene coded region of the envelope of hepatitis B virus employing the peptide immunogen of claim 1, said immunoassays being performed on serum taken from said human at different times.

1	150. A method for detecting the presence of
2	hepatitis B virus infection comprising effecting quantative
3 `	immunoassays on a serum sample taken from a human to
4	determine the amount of antibodies present therein which are
5	antibodies to an antigen coded by the pre-S gene coded
6	region of the envelope of the hepatitis B virus employing
7	the peptide immunogen of claim 1 and comparing the value
8	with a known standard.
9	151. A process for raising antibodies which
10	comprises introducing into an animal the peptide immunogen
11	of claim 1.
12	152. In a process for synthesizing His and Trp
13	containing peptides which includes the steps of
14	a. linking a first amino acid containing an
15	alpha-amino protecting group to a resin;
16	b. removal of said alpha-amino protecting group;
17	c. coupling a second amino acid containing an
18	alpha-amino protecting group to said first amino acid;
19	d. repeating steps b and c by coupling further
20	alpha-protected amino acids to produce a desired peptide
21	wherein at least one of said amino acids is His
22	and wherein at least one of said amino acids is Trp and the
23	e. cleaving the peptide from the resin and
24	removing remaining protective groups to said first amino
25	acids,
26	
27	wherein the improvement comprises substituting an
28	His(ImDNP) for said His, substituting a Trp(InFormyl) for
29	said Trp, removing said DNP prior to cleavage and said
- A	THROUGHUS OF DECEMBERING AVAIR SAN YAMANIDA CALA FORMUL

1	during said cleavage and said removing of protective group.
2	153. A hepatitis B vaccine comprising
3	a peptide containing an amino acid chain
4	corresponding to at least six consecutive amino acids within
5	the pre-S gene coded region of the envelope of HBV, said
6	vaccine free of an amino acid sequence corresponding to the
7	naturally occurring envelope proteins of hepatitis B virus,
8	and
9	a physiologically acceptable diluent.
10	154. A hepatitis B vaccine according to claim 153,
11	wherein said chain of amino acids is between sequence
12	position pre-S 120 and pre-S 174.
13	155. A hepatitis B vaccine according to claim 154,
14	wherein said chain of amino acids includes N-terminal
15	methionine at sequence position pre-S 120.
16	156. A hepatitis B vaccine according to claim 153,
17	wherein said chain of amino acids is between sequence
18	position pre-S 1 and pre-S 120.
19	157. A hepatitis B vaccine according to claim 153,
20	wherein said peptide contains a chain of at least 10 amino
21	acids.
22	158. A hepatitis B vaccine according to claim 153,
23	wherein said peptide contains a chain of at least 15 amino
24	acids.
25	159. A hepatitis B vaccine according to claim 153,
26	wherein said peptide contains a chain of at least 20 amino
27	acids.
28	160. A hepatitis B vaccine according to claim 153,
29	wherein said peptide contains a chain of at least 26 amino
30	MINETERI SOUR DEDUTES CONTOURS & CUSIN OF WE TEAST SO WILLIO

1	acids.
2	161. A hepatitis B vaccine according to claim 160,
3	wherein said chain is between and including sequence
4 .	positions pre-S 120 and pre-S 174.
5	162. A hepatitis B vaccine according to claim 161,
6	wherein said chain includes N-terminal methionine at
7	sequence position pre-S 120.
8	163. A hepatitis B vaccine according to claim 153,
9	wherein said chain is between and including sequence
10	position pre-S 15 and pre-S 120.
11	164. A hepatitis B vaccine according to claim 153,
12	wherein said chain is between and including sequence
13	position pre-S 15 and pre-S 55.
14	165. A hepatitis B vaccine according to claim 153,
15	wherein said chain is between and including sequence
16	position pre-S 90 and pre-S 120.
17	166. A hepatitis B vaccine according to claim 153,
18	wherein said chain is between and including sequence
19 20	position pre-S 10 and pre-S 40.
21	167. A hepatitis B vaccine according to claim 153,
22	wherein said chain corresponds to amino acids in the ayw
23	subtype of the pre-S region.
24	168. A hepatitis B vaccine according to claim 153,
25	wherein said chain corresponds to amino acids in the adyw
26	subtype of the pre-S region.
27	169. A hepatitis B vaccine according to claim 153,
28	wherein said chain corresponds to amino acids in the adw2
29	subtype of the pre-S region.

1	. 170. A hepatitis B vaccine according to claim 153,
2	wherein said chain corresponds to amino acid in the adw
3	subtype of the pre-S region.
4	171. A hepatitis B vaccine according to claim 153,
5	wherein said chain corresponds to amino acids in the adr
6	subtype of the pre-S region.
7	172. A hepatitis B vaccine according to claim 153,
8	wherein said vaccine is free of any serum proteins.
9	173. A heptatis B vaccine according to claim 153,
10	wherein said chain is MQWNSTAFHQTLQDPRVRGLYLPAGG.
11	
12	174. A hepatitis B vaccine according to claim 153,
13	wherein said chain is MGTNLSVPNPLGFFPDHQLDP.
14	175. A hepatitis B vaccine according to claim 153,
15 -	wherein said chain is PAFGANSNNPDWFNPVKDDWP.
16	176. A hepatitis B vaccine according to claim 153,
17	wherein said chain is PQAMQWNSTAFHQTLQDP.
18	177. A hepatitis B vaccine according to claim
19	153, wherein said chain is PASTNRQSGRQPTPISPPLRDSHP.
20	178. A hepatitis B vaccine according to claim 153,
21	wherein said chain is PAPNIASHISSISARTGDP.
22	179. A hepatitis B vaccine according to claim
23	153, wherein said chain is MGGWSSKPRKGMGTNLSVPNP.
24	180. A hepatitis B vaccine according to claim 153,
25	wherein said chain is PAFGANSNNPDWDFNPVKDDWP.
26	181. A hepatitis B vaccine according to claim 153,
27	wherein said chain is QVGVGAFGPRLTPPHGG.
28	182. A hepatitis B vaccine according to claim 153,
:9	wherein said chain is MGGWSSKPRKG.

1	183. A hepatitis B vaccine according to claim 153,
2	wherein said chain is MGGWSSKPRQG.
3 .	184. A hepatitis B vaccine according to claim
4	153, wherein said vaccine is free of an amino acid sequence
5	corresponding to the entire S gene coded region of the env
6	gene product of hepatitis B virus.
7	185. A hepatitis B vaccine according to claim
8	153, wherein said peptide has no more than 100 amino acids.
9	186. A hepatitis B vaccine according to claim
10	153, wherein said peptide has no more than 40 amino acids.
11	187. A hepatitis B vaccine according to claim
12	153, wherein said peptide has no more than 30 amino acids.
13.	188. A hepatitis B vaccine according to claim
14	153, wherein said peptide is capable of forming neutralizing
15	antibodies to hepatitis B virus in a humans.
16	189. A hepatitis B vaccine according to claim 153,
17	wherein said peptide is linked to a carrier.
18	190. A hepatitis B vaccine according to claim 189,
19	wherein said peptide is covalently linked to a carrier.
20	
21	191. A hepatitis B vaccine according to claim
22	190, wherein said peptide is covalently linked to a lipid
23	vesicle carrier.
24	192. A hepatitis B vaccine according to claim 191,
25	wherein said lipid vesicle is stabilized by cross-linking.
26	193. A hepatitis B vaccine according to claim
27	153, wherein said chain is pre-S(120-145).
28	194. A hepatitis B vaccine according to claim
29	153, wherein said chain is pre-S(12-32).

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195. A hepatitis B vaccine accord 1 2 173, wherein said chain is pre-S(117-134). 3 196. A hepatitis B vaccine according to claim 4 153, wherein said chain is pre-S(94-117).

> 197. A hepatitis B vaccine according to claim 153, wherein said chain is pre-S(153-171).

> 198. A hepatitis B vaccine according to claim 153, wherein said chain is pre-S(1-21).

> 199. A hepatitis B vaccine according to claim 153, wherein said chain is pre-S(57-73).

200. A method of protecting a human against becoming infected with hepatitis B comprising administering to said human an effective dosage of a vaccine according to claim 153.

201. A method for detecting the presence of hepatitis B virus infection comprising effecting quantative immunoassays on a serum sample taken from a human to determine the amount of antigens coded by the pre-S gene coded region of the envelope of the hepatitis B virus employing antibodies to the peptide immunogen of claim 1 and comparing the value with a known standard.

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FIG. I

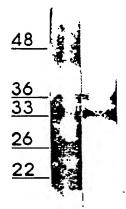
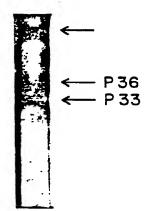


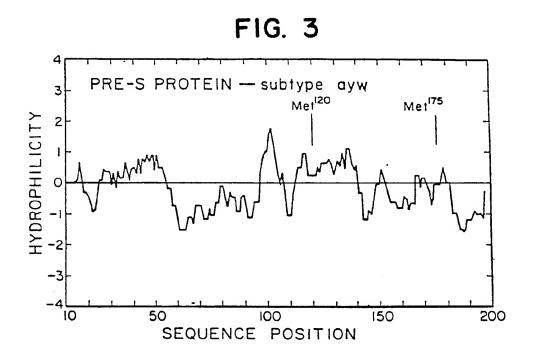
FIG. 6



# F16. 2

Pre-S protein

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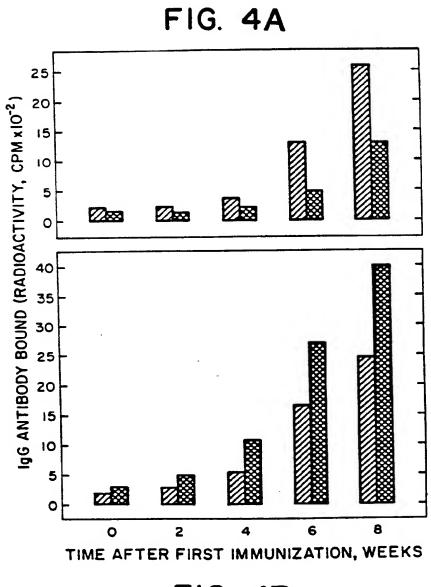
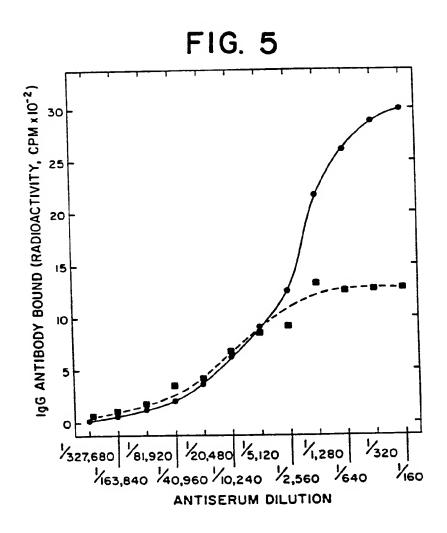
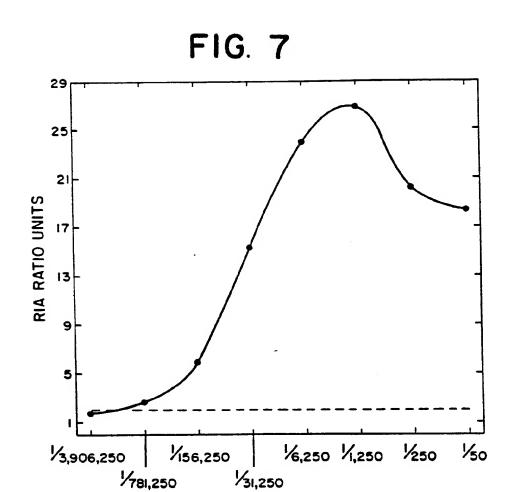
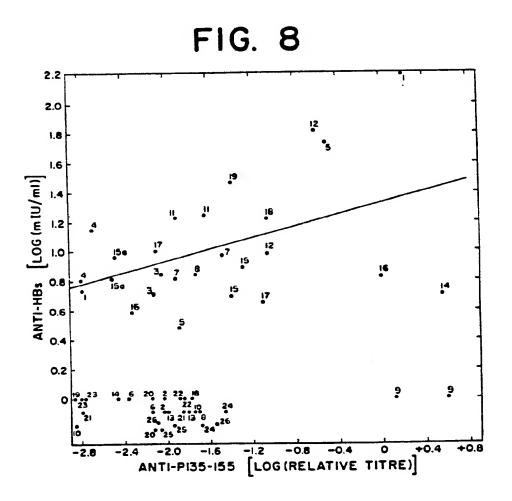


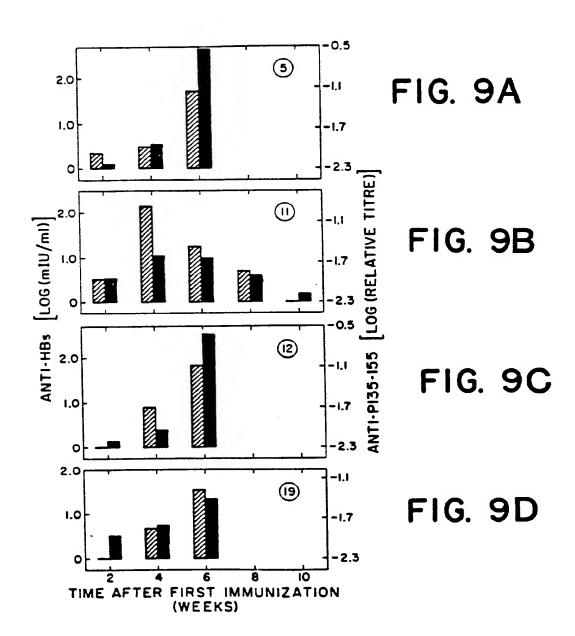
FIG. 4B

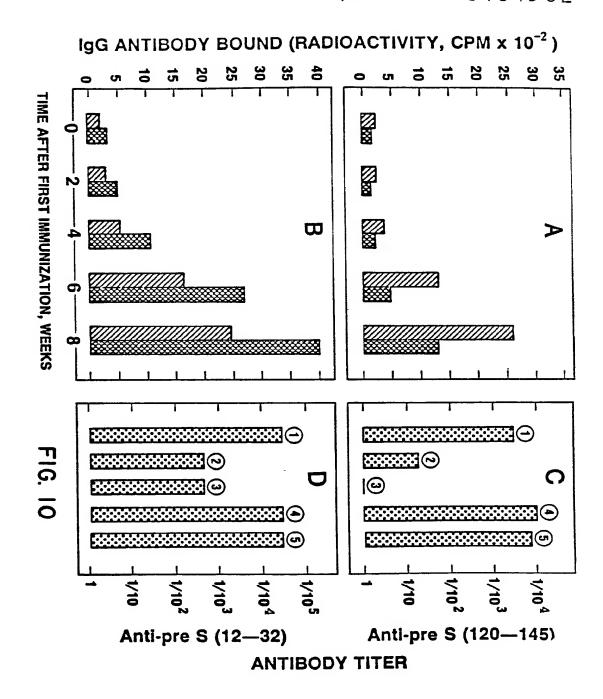


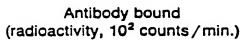


DILUTION OF HBsAg - POSITIVE SERUM









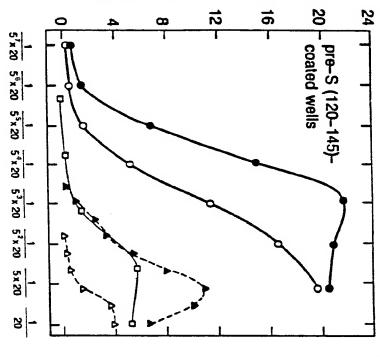
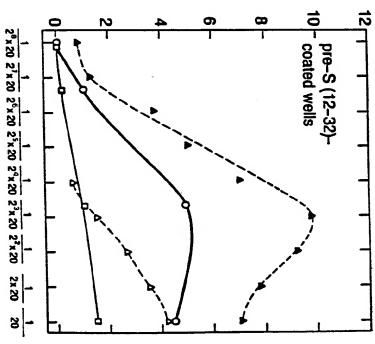
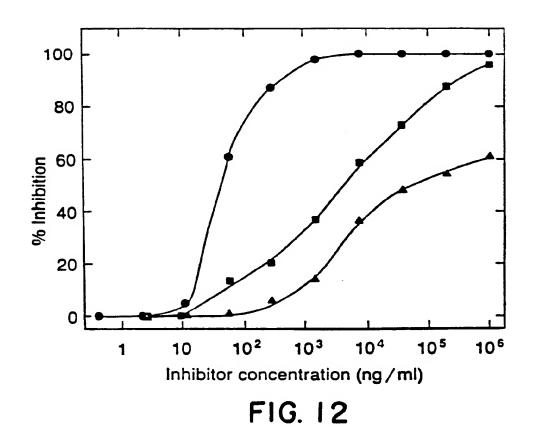


FIG. II

Serum dilution



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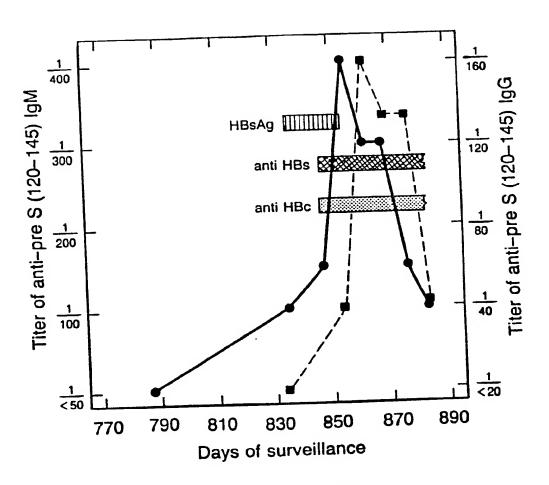


FIG. 13

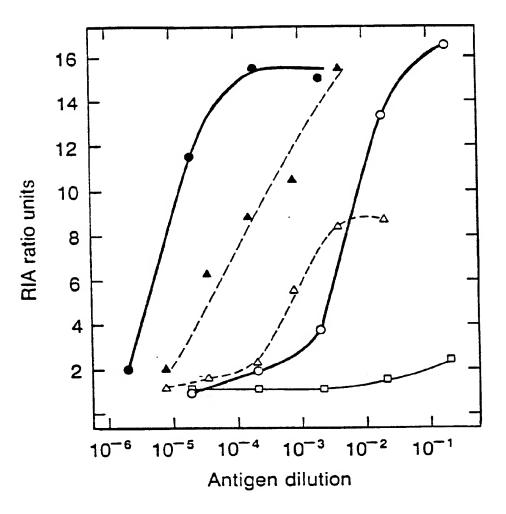
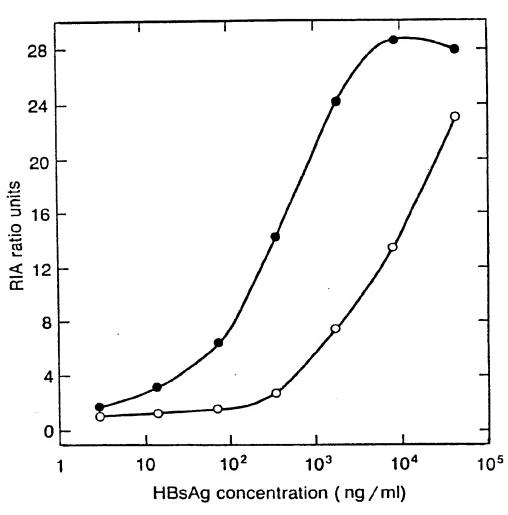


FIG. 14



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FIG. 15